



Mitochondria in Space: The Data in Nutshell

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President

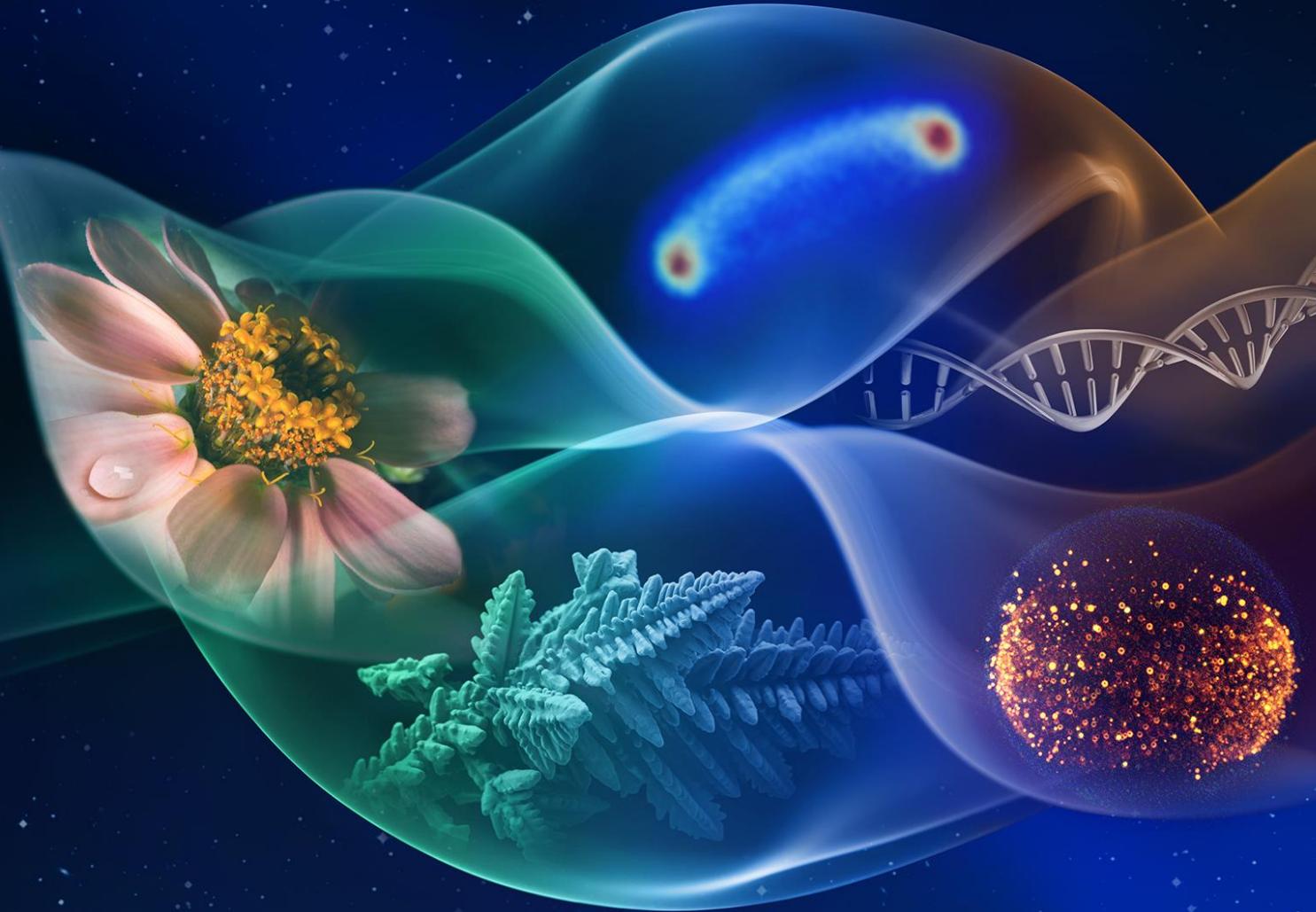
COVID-19 International Research Team (COV-IRT)

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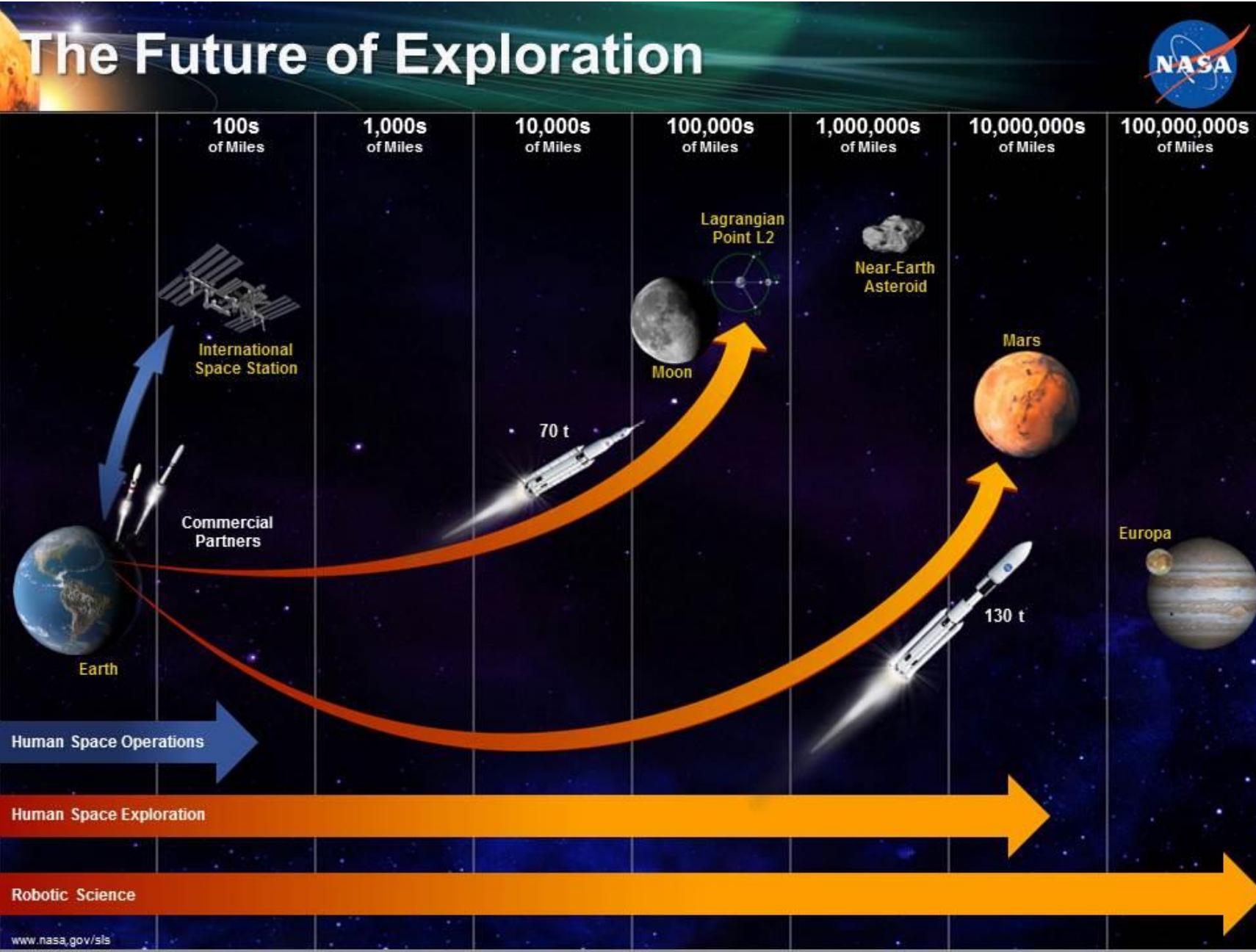
www.cov-irt.org



Background on Spaceflight, Experiments, and Resources for the Data That was used for the Mitochondrial project

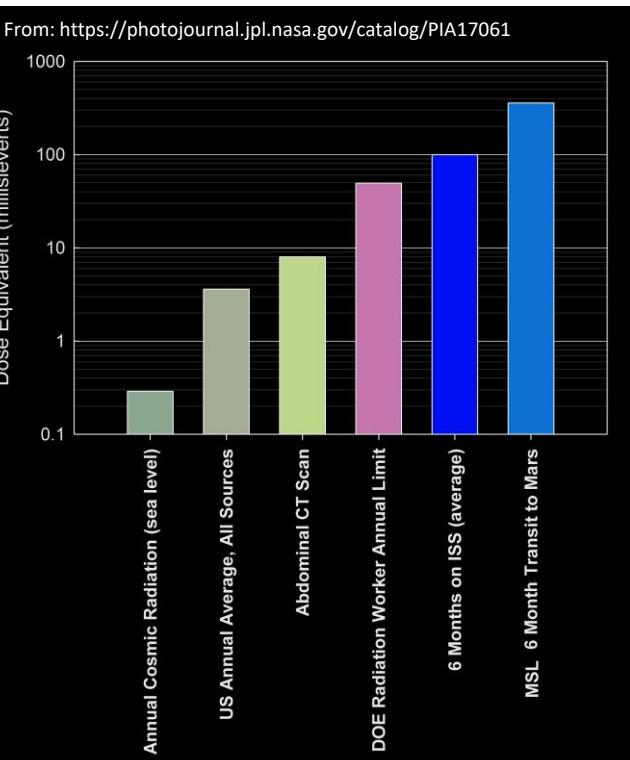
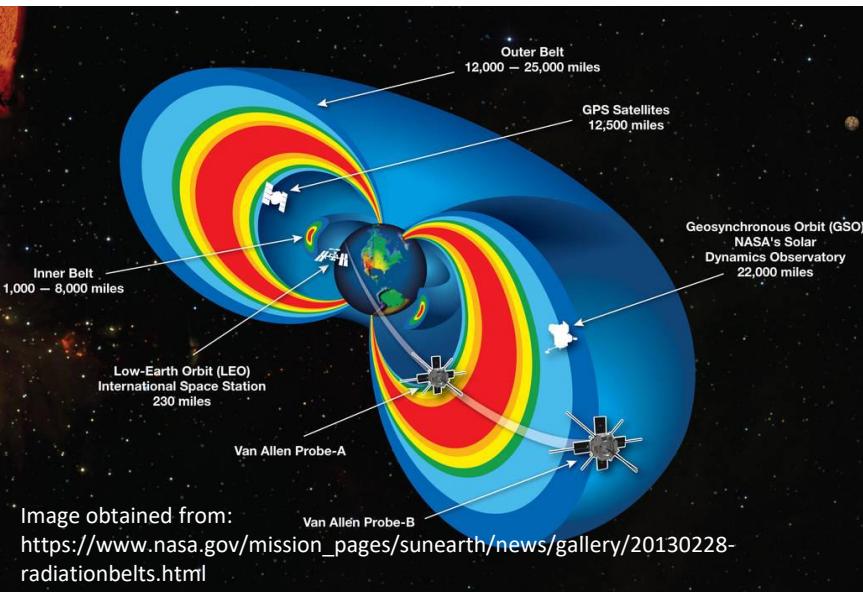
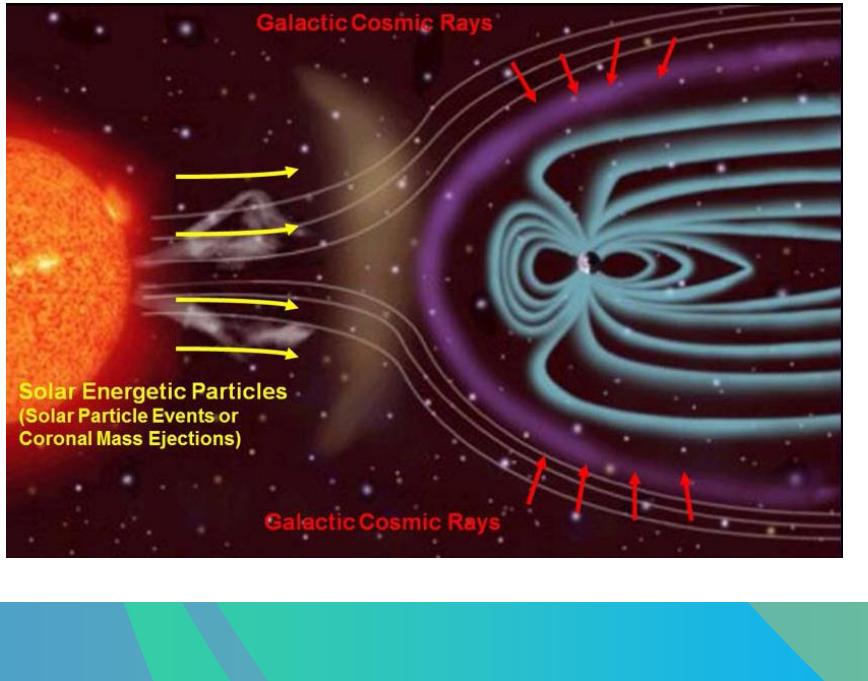


Why Care About Space Biology Research



Space Environment

Radiation



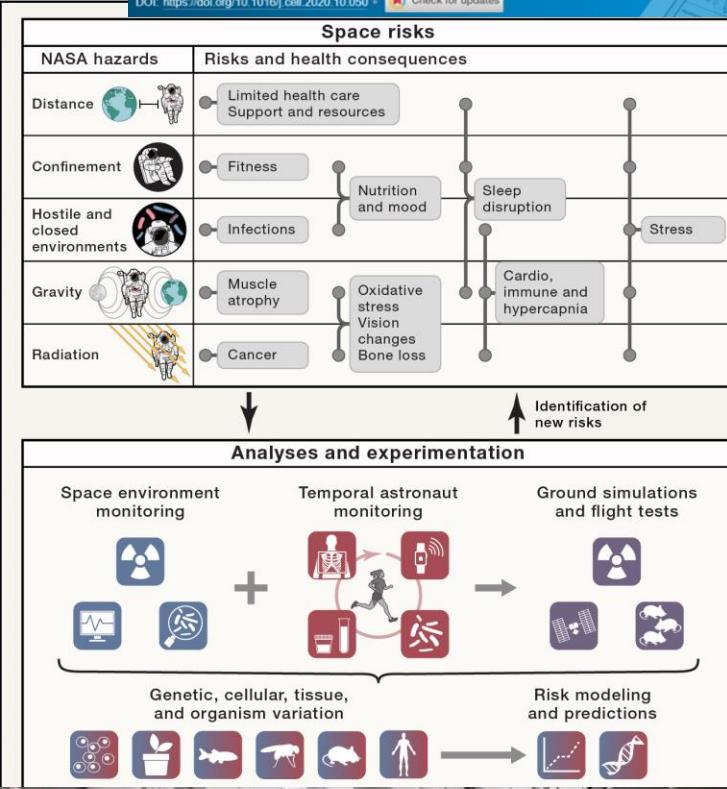
Microgravity



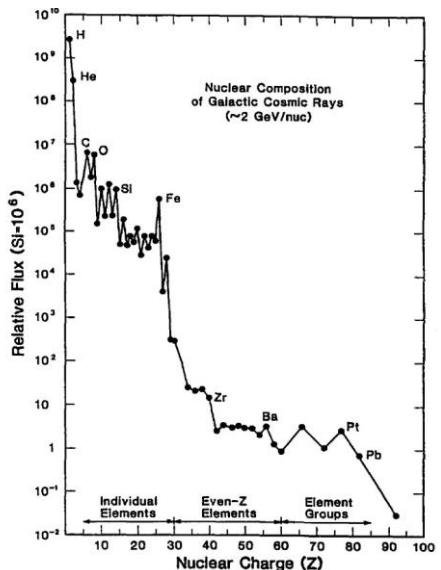
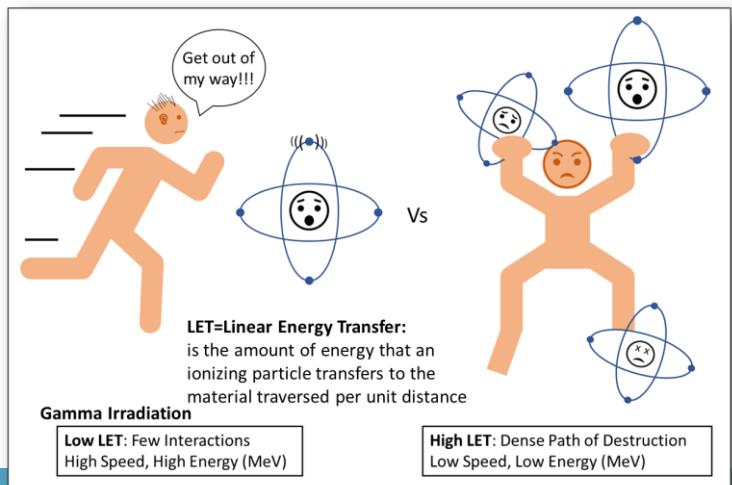
Cell
Fundamental Biological Features of Spaceflight: Advancing the Field to Enable Deep-Space Exploration

Ebrahim Afshinnekoo ³⁴ • Ryan T. Scott ³⁴ • Matthew J. MacKay ³⁴ • ... Sylvain V. Costes ³⁵ • ... Christopher E. Mason ³⁵ • Afshin Beheshti ³⁵ • Show all authors • Show footnotes

DOI: <https://doi.org/10.1016/j.cell.2020.10.050> • Check for updates



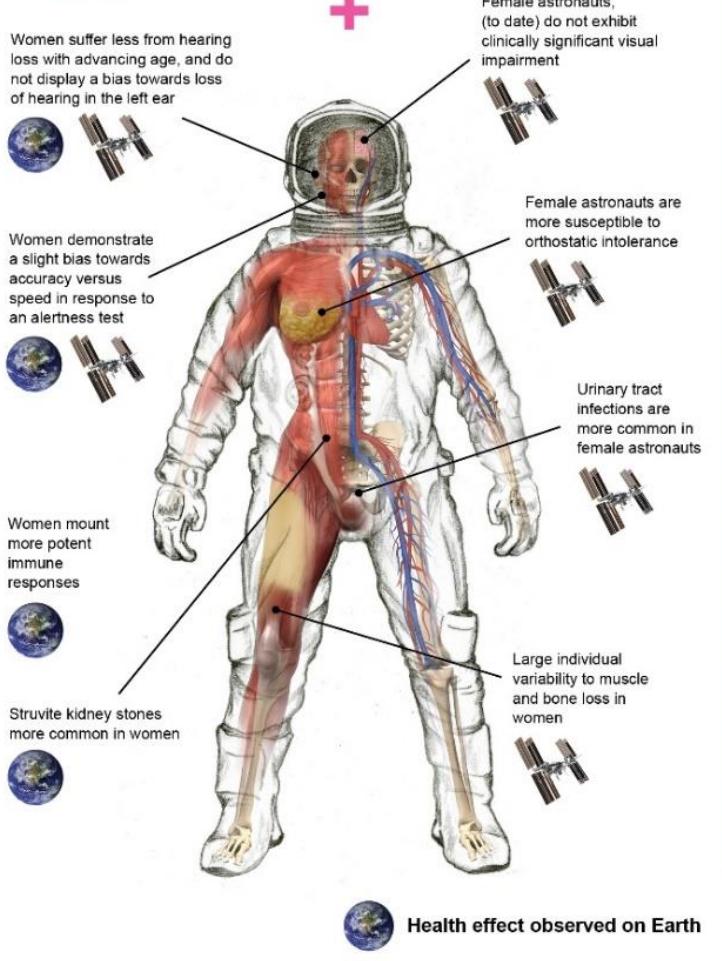
Galactic Cosmic Radiation (High LET) – a cautionary tale



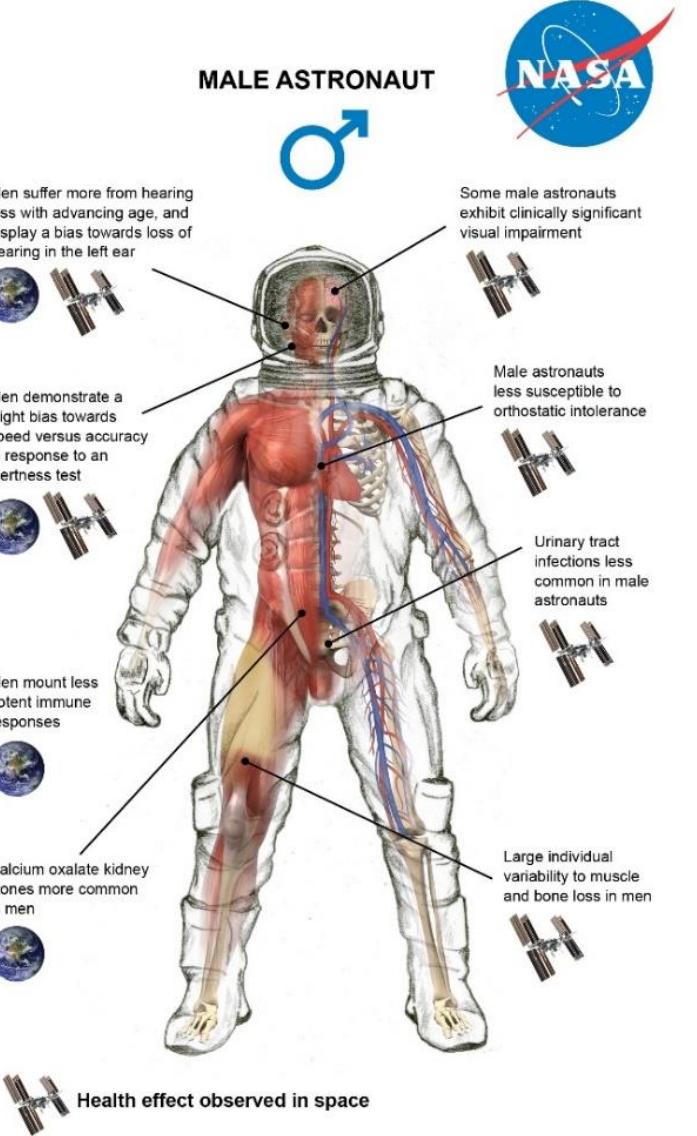
Space Environment Health Risks On Astronauts



FEMALE ASTRONAUT



MALE ASTRONAUT



Mission and long-term health risks							
Mission type	Low Earth orbit	Low Earth orbit	Deep space sortie	Lunar visit/habitat	Deep space journey	Deep Space	Planetary visit/habitat
Mission duration	6 months	12 months	1 month	12 months	12 months	Weeks/months	Months
Return duration	<= 1 day	<= 1 day	< 5 days	5 days	12 months	Weeks/months	Months
Radiation	Van Allen	Van Allen	Deep Space	Lunar	Deep Space	Variable	Variable
Gravity	Micro	Micro	Micro	Micro	1/6g	Micro	Variable
Health risks	Mission	Long-term	Mission	Long-term	Mission	Long-term	Mission
Renal	●	●	●	●	●	●	●
Medical	●	●	●	●	●	●	●
SANS	●	●	●	●	●	●	●
Arrhythmia	●	●	●	●	●	●	●
BMed	●	●	●	●	●	●	●
Occupant protection	●	●	●	●	●	●	●
Hypobaric hypoxia	●	●	●	●	●	●	●
EVA	●	●	●	●	●	●	●
Degen	●	●	●	●	●	●	●
CNS	●	●	●	●	●	●	●
Team	●	●	●	●	●	●	●
Sleep	●	●	●	●	●	●	●
Sensorimotor	●	●	●	●	●	●	●
Cancer	●	●	●	●	●	●	●
Muscle	●	●	●	●	●	●	●
Aerobic	●	●	●	●	●	●	●
Immune	●	●	●	●	●	●	●
Microhost	●	●	●	●	●	●	●
DCS	●	●	●	●	●	●	●
Stability	●	●	●	●	●	●	●
OI	●	●	●	●	●	●	●
ARS	●	●	●	●	●	●	●
Dust	●	●	●	●	●	●	●

Planned human missions



Type of Experiments Related to Space Biology

Experiments Done in Space

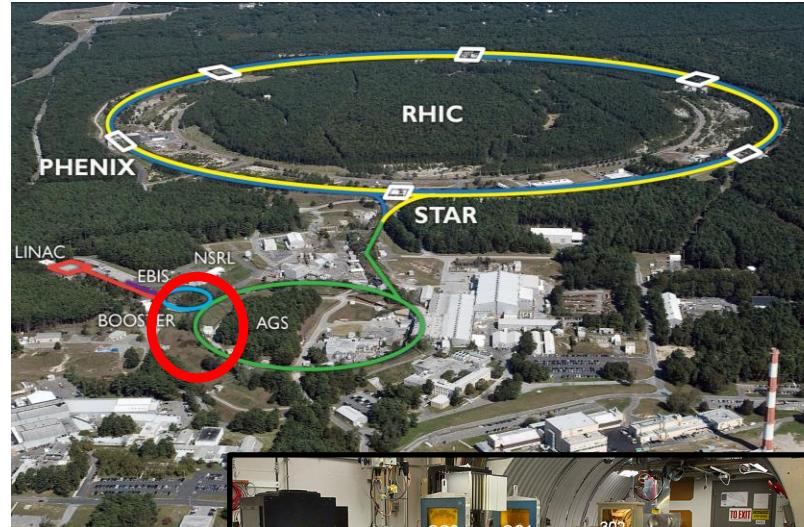


FLIGHT HABITAT, DAY 2 (DARK CYCLE)



Space Radiation Simulated Experiments

Brookhaven National Laboratory



jove

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ABSTRACT

INTRODUCTION

PROTOCOL

RE

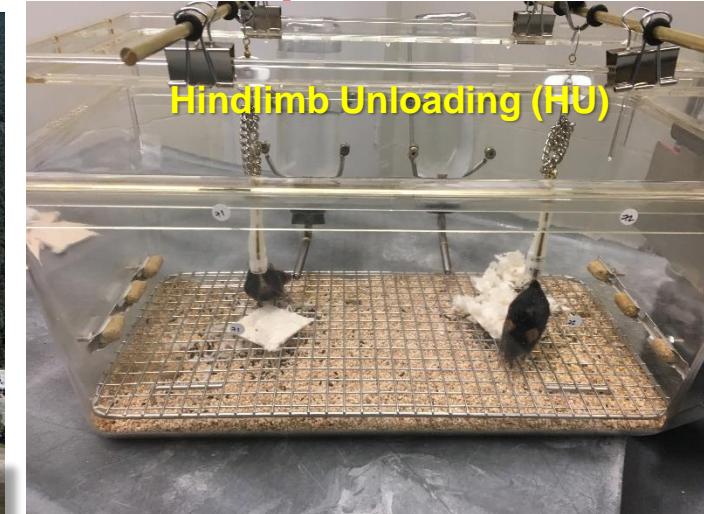
GENETICS

Exploring the Effects of Spaceflight on Mouse Physiology using the Open Access NASA GeneLab Platform

Afshin Beheshti¹, Yasaman Shirazi-Fard², Sungjin Choi¹, Daniel Berrios³, Samrawit G. Gebre¹, Jonathan M. Galazka², Sylvain V. Costes²

¹WYLE Labs, Space Biosciences Division, NASA Ames Research Center, ²Space Biosciences Division, NASA Ames Research Center, ³USRA, NASA Ames Research Center

Microgravity Simulated Experiments



Partial Weight Bearing Rat Model



Seward Rutkove



Marie Mortreux



Beth Israel Deaconess
Medical Center

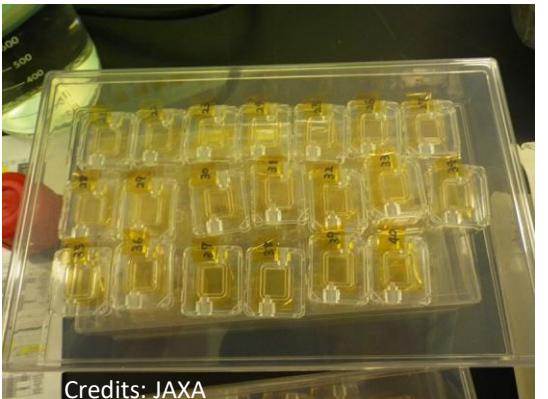
A teaching hospital of
Harvard Medical School

<https://www.rutkovelab.org/nasa/>

Other Types of Experiments on the ISS

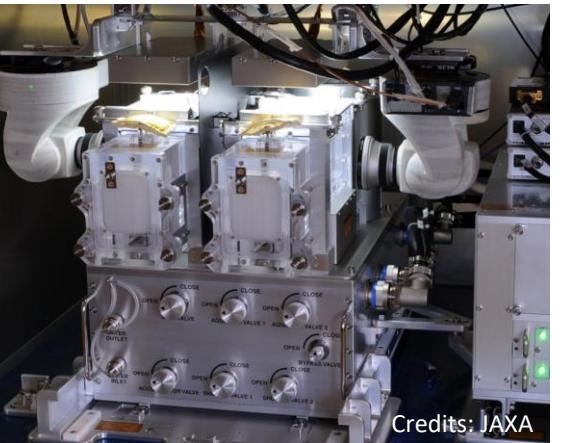


Credits: NASA

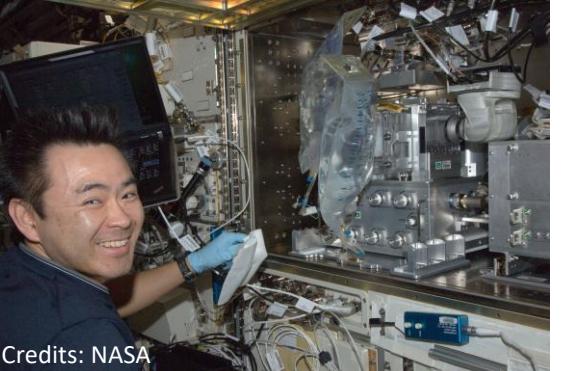


Credits: JAXA

C. elegans culture chambers for the Space Aging experiment aboard the ISS



Credits: JAXA

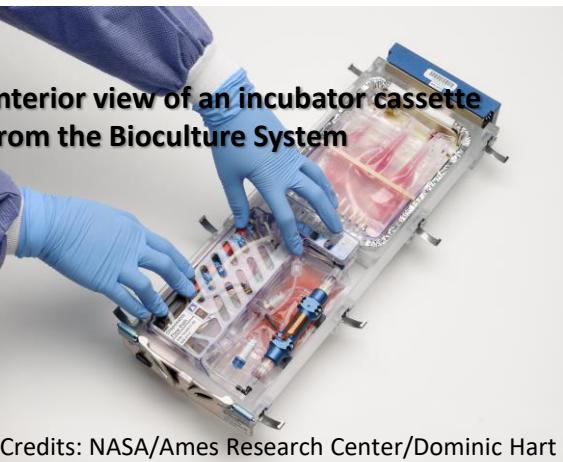


Credits: NASA

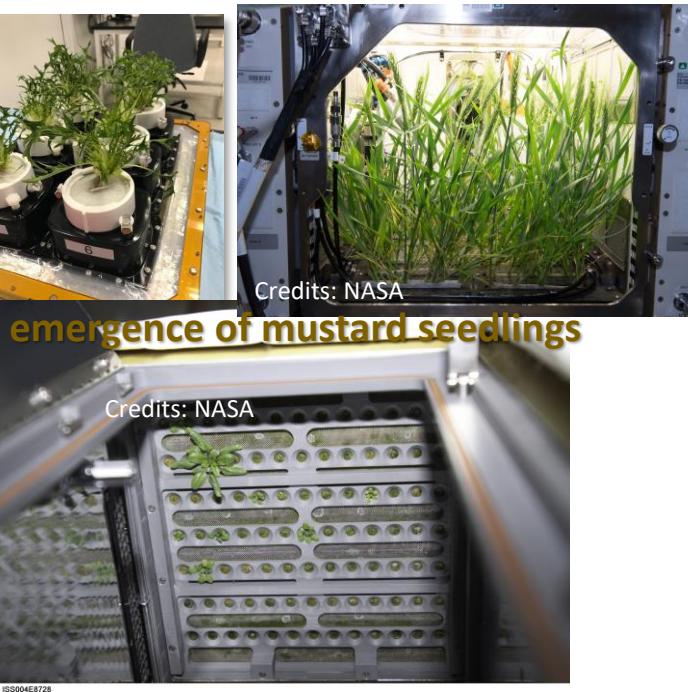
The Zebrafish Muscle investigation employs the ISS Aquatic Habitat, an aquarium in microgravity.



Credits: NASA

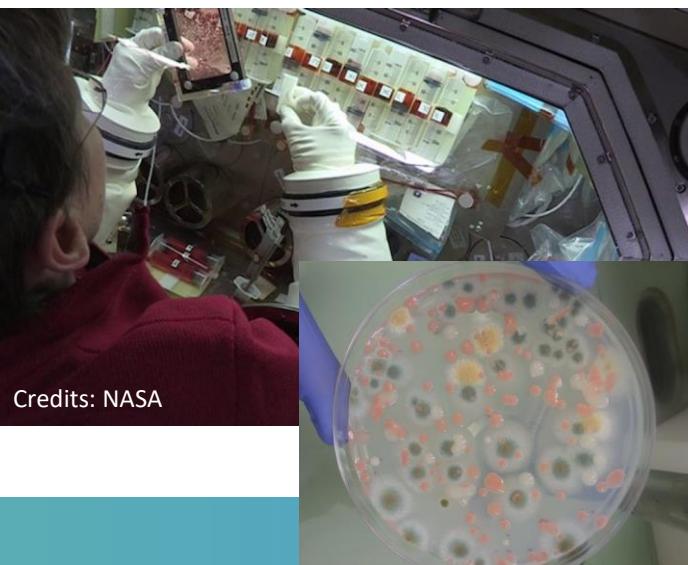


Credits: NASA/Ames Research Center/Dominic Hart



Credits: NASA

ISS004E8728



Credits: NASA

Lots of Papers Published on Space Biology in 11/2020!!

<https://www.cell.com/c/the-biology-of-spaceflight>



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Related resources ↓ Contributing lab groups ↓ Consortia Hub ↗

The biology of spaceflight

As humankind reaches for the stars to journey to the next frontier in space, research on spaceflight biology is critical for understanding how living systems, including human health, may be affected by spaceflight and space exploration. This special collection on the biology of spaceflight, published in *Cell* and other Cell Press journals, includes research articles, short communications, and a review article that cover studies with model systems and astronaut samples. The work, which was done in collaboration between NASA and other space agencies around the world, uncovers the impact of known hazards of spaceflight, such as radiation and microgravity, and discusses the standards for multi-omics from space and the preparations needed for Mars and other missions in the next two decades.



Log in



ARTICLE | VOLUME 183, ISSUE 5, P1185-1201.E20, NOVEMBER 25, 2020

Comprehensive Multi-omics Analysis Reveals Mitochondrial Stress as a Central Biological Hub for Spaceflight Impact

Willian A. da Silveira ²³ • Hossein Fazelia ²³ • Sara Brin Rosenthal ²³ • ...

Christopher E. Mason ²⁴ • Sylvain V. Costes ²⁴ • Afshin Beheshti ^{24, 25} Show all authors

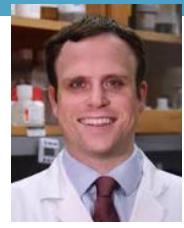
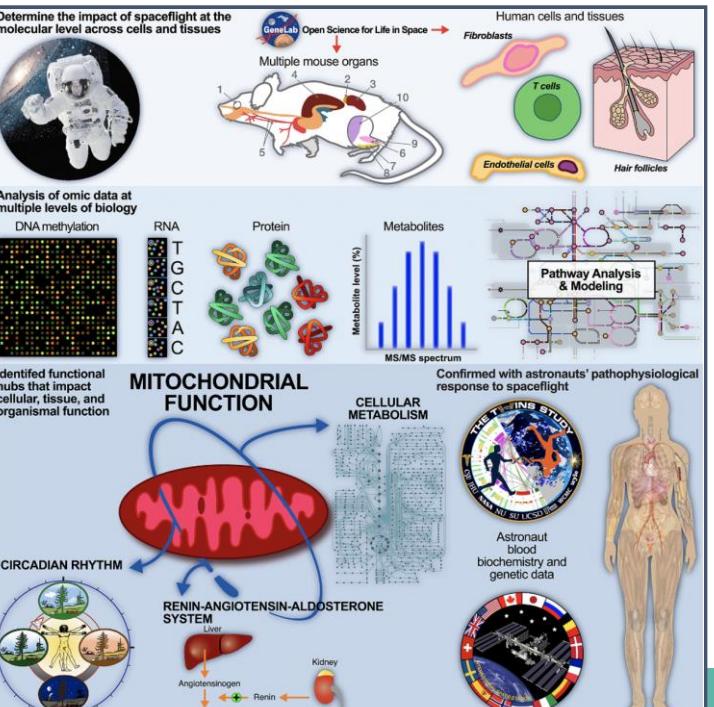
Show footnotes DOI: <https://doi.org/10.1016/j.cell.2020.11.002>

PlumX Metrics



Highlights

- Multi-omics analysis and techniques with NASA's GeneLab platform
- The largest cohort of astronaut data to date utilized for analysis
- Mitochondrial dysregulation driving spaceflight health risks
- NASA Twin Study data validates mitochondrial dysfunction during space missions



GeneLab (genelab.nasa.gov)



Open Science for Life in Space



Open Science for Life in Space

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Keywords



Welcome to NASA GeneLab - the first comprehensive space-related omics database; users can upload, download, share, store, and analyze spaceflight and spaceflight-relevant data from experiments using model organisms.



Data Repository

Search and upload spaceflight datasets



Analyze Data

Perform large-scale analysis of biological omics data



Environmental Data

Radiation data collected during experiments conducted in space



Collaborative Workspace

Share, organize and store files



Submit Data BETA

Have space-relevant data to submit to GeneLab?



Visualize Data

Interact with GeneLab processed data

Members are now group leads

AWG Members Per Group:

Animal

39

Multi-Omics/System Biology

250+

Plants

44

Microbes

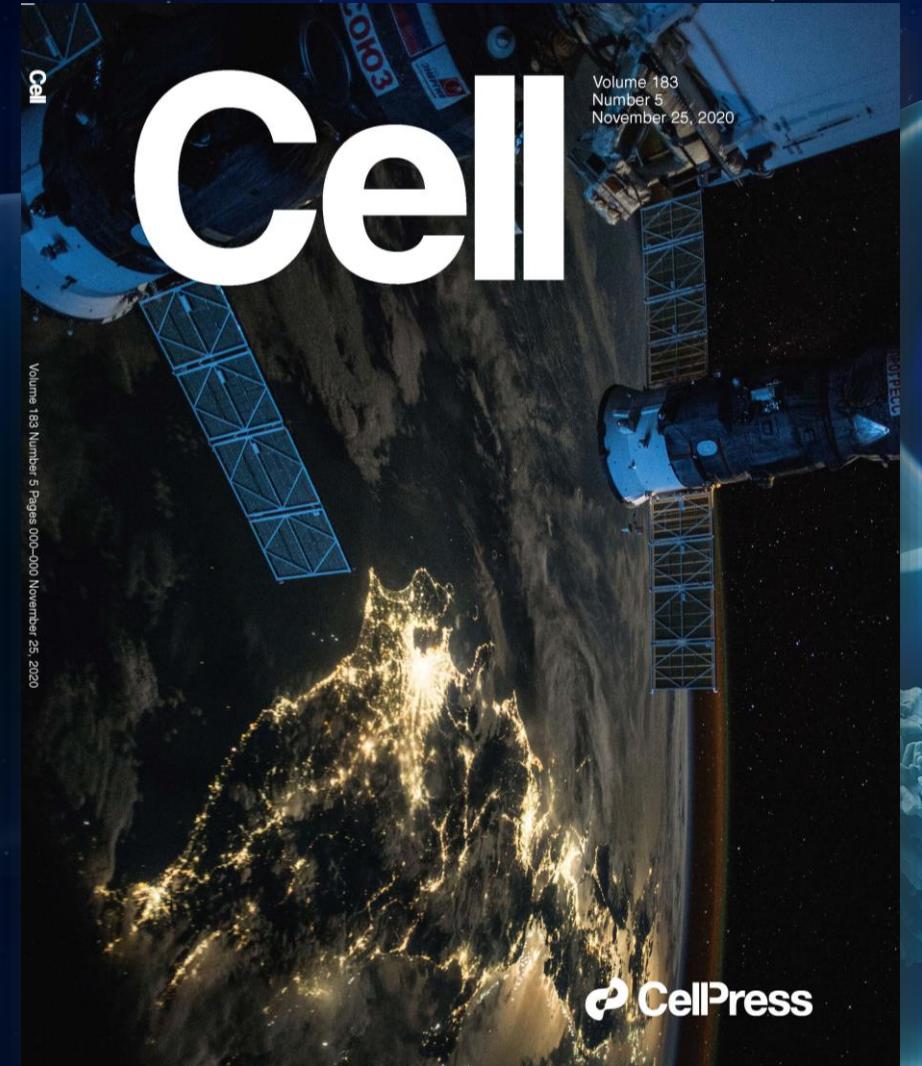
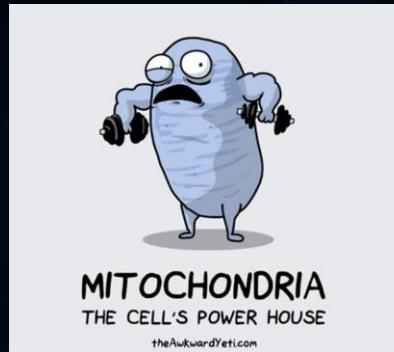
25

I lead the Multi-Omics Group

GeneLab Analysis Working Groups (AWGs) consist of 300+ scientists from multiple space agencies, international institutions, and industry. Scientists meet monthly with each group to analyze data in the GeneLab repository. Majority of members are non-NASA PI's – many have applied for NASA funding following AWG interactions.

<https://genelab.nasa.gov/awg/join>

Spaceflight Impact on the Mitochondria



Cell

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ARTICLE | VOLUME 183, ISSUE 5, P1185-1201.E20, NOVEMBER 25, 2020

Comprehensive Multi-omics Analysis Reveals Mitochondrial Stress as a Central Biological Hub for Spaceflight Impact

Willian A. da Silveira ²³ • Hossein Fazelia ²³ • Sara Brin Rosenthal ²³ • ...
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Show footnotes DOI: <https://doi.org/10.1016/j.cell.2020.11.002>

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- Multi-omics analysis and techniques with NASA's GeneLab platform
- The largest cohort of astronaut data to date utilized for analysis
- Mitochondrial dysregulation driving spaceflight health risks
- NASA Twin Study data validates mitochondrial dysfunction during space missions

Determine the impact of spaceflight at the molecular level across cells and tissues

GeneLab Open Science for Life in Space → Human cells and tissues
Multiple mouse organs Fibroblasts
T cells
Endothelial cells Hair follicles

Analysis of omic data at multiple levels of biology

DNA methylation RNA Protein Metabolites

MS/MS spectrum

Identified functional hubs that impact cellular, tissue, and organismal function

Pathway Analysis & Modeling

Confirmed with astronauts' pathophysiological response to spaceflight

THE TWIN STUDY Astronaut blood biochemistry and genetic data

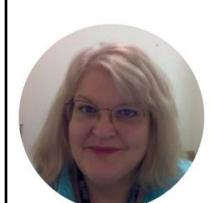
MITOCHONDRIAL FUNCTION
CIRCADIAN RHYTHM
RENIN-ANGIOTENSIN-ALDOSTERONE SYSTEM
CELLULAR METABOLISM
Liver Kidney

Angiotensinogen Angiotensin I Renin

Analysis Working Group (AWG) Members and Others Involved



Willian da Silveira



Deanne Taylor



Hossein Fazelinia



Komal Rathi



Douglas Wallace



Larry Singh



Benjamin
Stear



"Jimmy" Man S
Kim



Kathleen Fisch



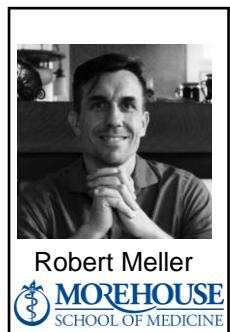
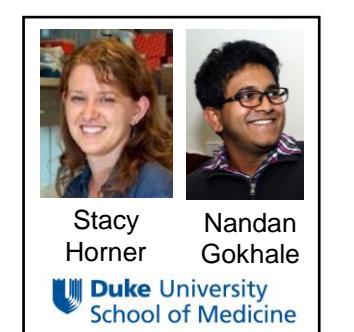
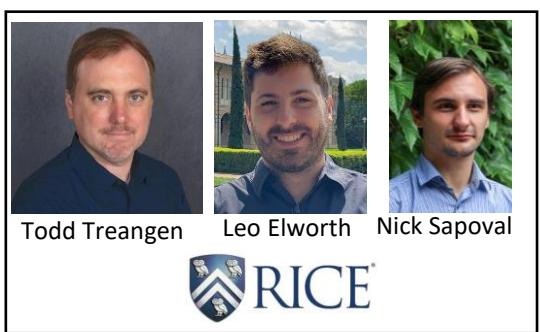
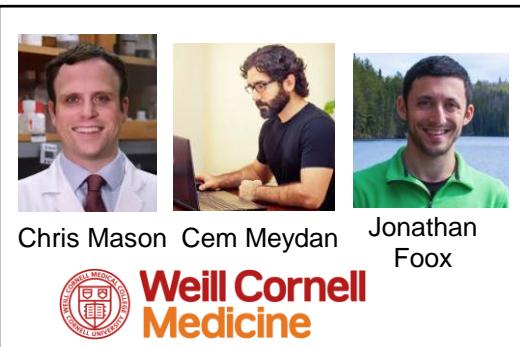
UNIVERSITY OF CALIFORNIA, SAN DIEGO
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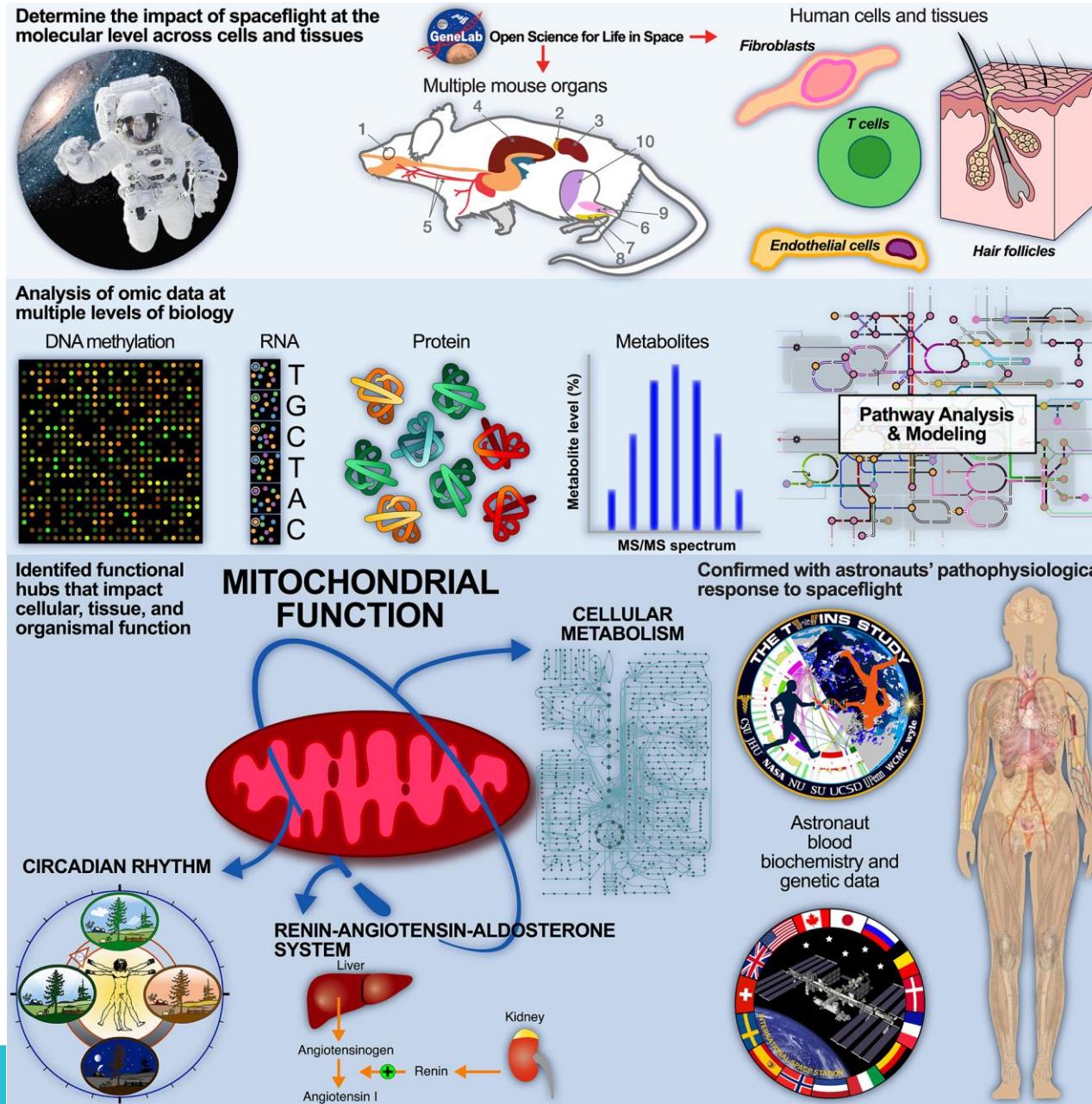
Brin Rosenthal



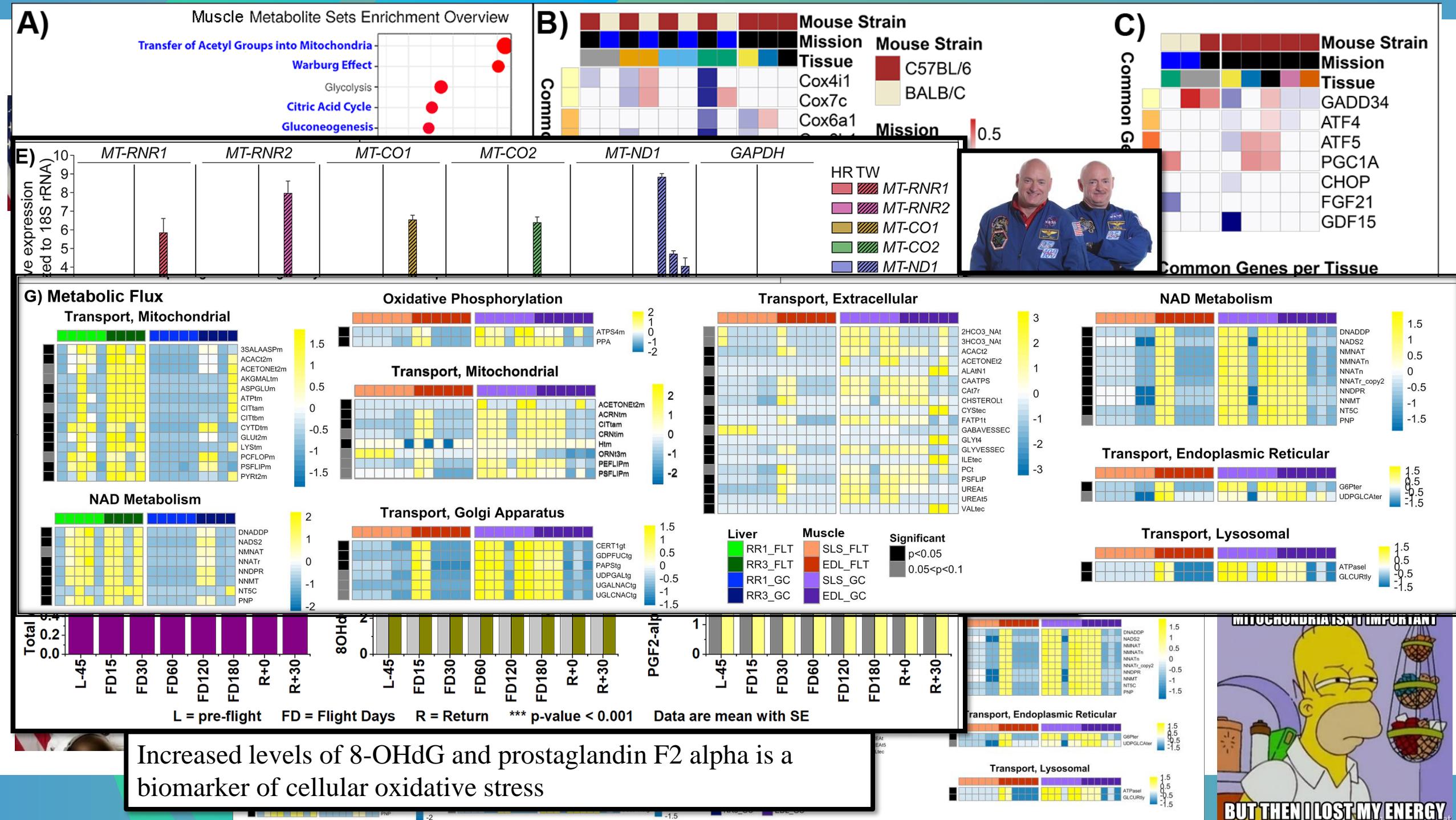
Jonathan
Schisler



Overview of Data Utilized for this Work and Project



Largest amount of astronaut data in one paper!!



Ongoing work: Detailed Mitochondrial Analysis on Data from Astronauts

OSD-530 Version 1

Cell-free RNA analysis of plasma samples collected from six astronauts in JAXA Cell-Free Epigenome (CFE) Study

Submitted Date: 13-Sep-2022
Initial Release Date: 01-Dec-2022

Study

GeneLab ID: GLDS-530
DOI: 10.26030/r2xr-h731

Cite this Study

6 astronauts 120 days on the ISS

A) OXPHOS Complex I

B) OXPHOS Complex II

C) OXPHOS Complex III

D) OXPHOS Complex IV

E) mtDNA OXPHOS Complexes

F) OXPHOS Complex V

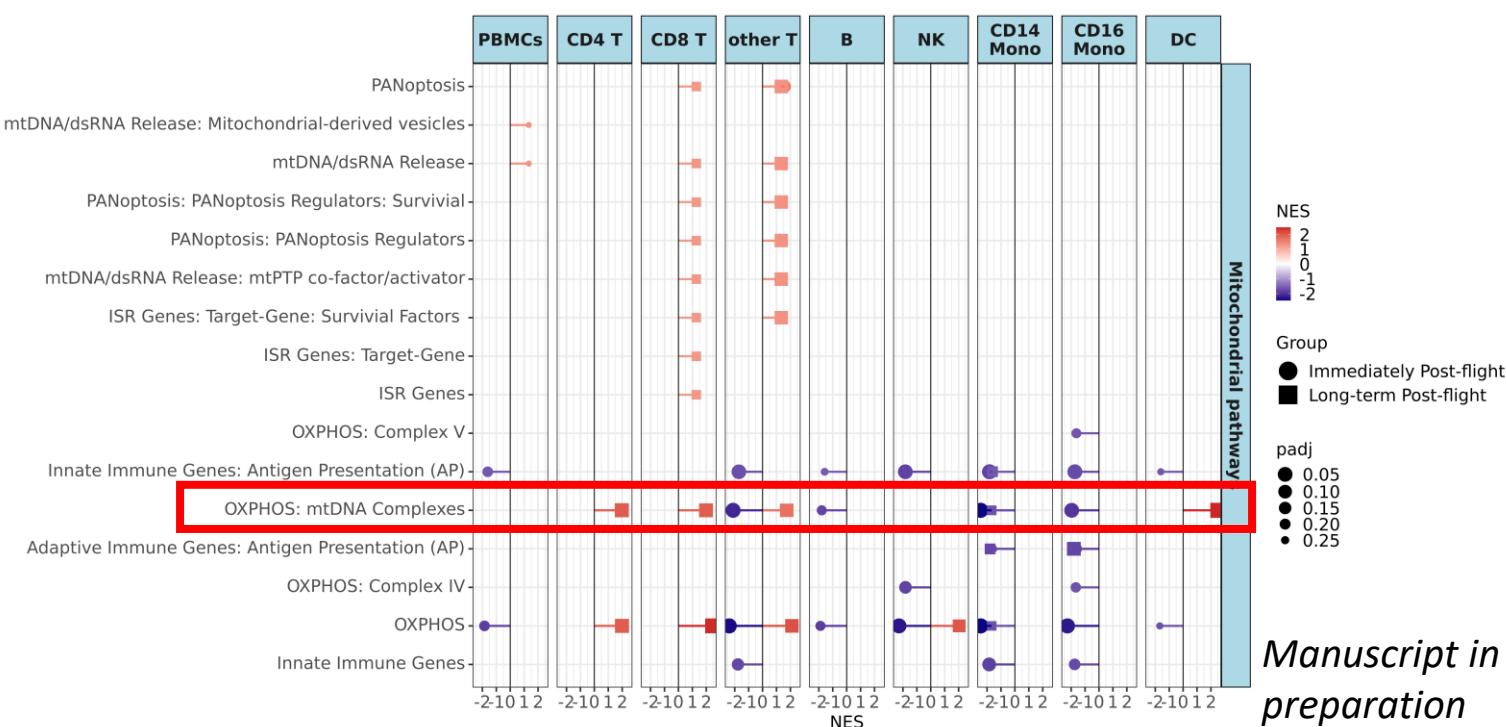
G) Flight vs Pre-Flight

H) Post-Flight vs Pre-Flight

I) Post-Flight vs Flight



4 astronauts 3 days 590km in elevation (accumulated radiation dose equivalent to ~9 months on ISS)



**Can we mitigate
the damage
caused to the
mitochondria in
space?
Short answer:
Maybe with
miRNAs!!**



ARTICLE | ONLINE NOW, 108448

Circulating miRNA Spaceflight Signature Reveals Targets for Countermeasure Development

Sherina Malkani ²² • Christopher R. Chin ²² • Egle Cekanaviciute ²² • ... Peter Graham •

Christopher E. Mason ²³ Afshin Beheshti ^{23, 24} Show all authors • Show footnotes

Open Access • Published: November 25, 2020 • DOI: <https://doi.org/10.1016/j.celrep.2020.108448>

PlumX Metrics



Highlights

- Spaceflight miRNA signature validated in multiple organism models
- Components of miRNA signature related to space radiation and microgravity
- Downstream targets and circulating dependence of miRNAs in NASA Twins Study
- Inhibition of key microvasculature miRNAs mitigates space radiation impact

OBJECTIVE
Identify circulating miRNAs impacted by spaceflight

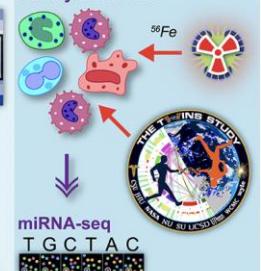


RESULTS
miRNA signature from multiple rodent models

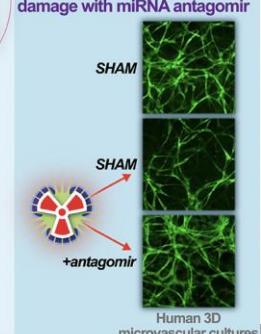


VALIDATION IN HUMAN CELLS AND ASTRONAUTS

Primary blood cells

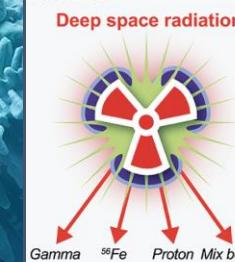


Rescue radiation-dependent damage with miRNA antagonist



MODELS

Deep space radiation



Simulated microgravity

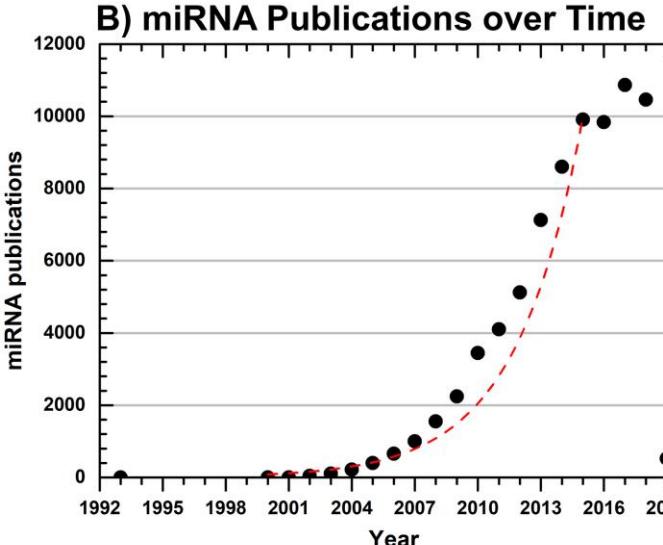
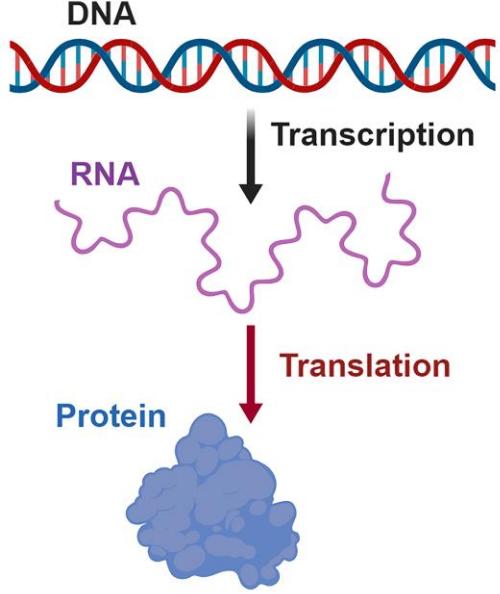


Pathways and diseases



What are miRNAs and why study miRNAs?

A) Classical View of Molecular Biology



C) New Understanding of Molecular Biology

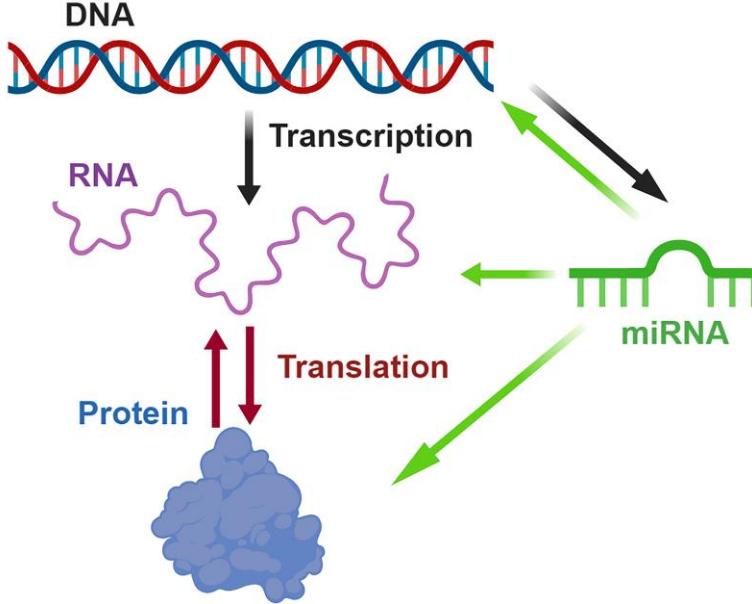
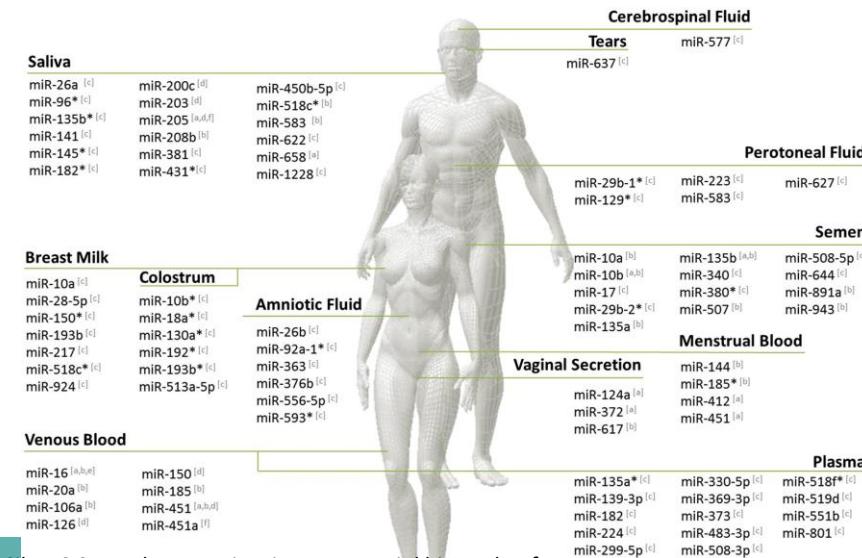


Figure from Vanderburg and Beheshti, MicroRNAs (miRNAs), the Final Frontier: The Hidden Master Regulators Impacting Biological Response in All Organisms Due to Spaceflight, THREE, 2020. (https://three.jsc.nasa.gov/articles/miRNA_Beheshti.cfm)

- A single miRNA has been estimated to regulate 100s to 1000s of mRNAs.
- miRNAs are ~22nt
- Due to the size and stability of the miRNAs, it can float freely in the blood. →
- miRNAs are now known to be involved in all aspects of diseases.
- miRNAs are not only found in mammals, but everything else living: plants, microbes, fish, C. Elegans, fruit flies, insects, etc...
- miRNAs are highly conserved across species.



Mitochondrial miRNAs (mitomiRs) Related to Countermeasures

Cell. Mol. Life Sci. (2017) 74:631–646
DOI 10.1007/s0018-016-2342-7

Cellular and Molecular Life Sciences



Life Sciences

Volume 164, 1 November 2016, Pages 60–70



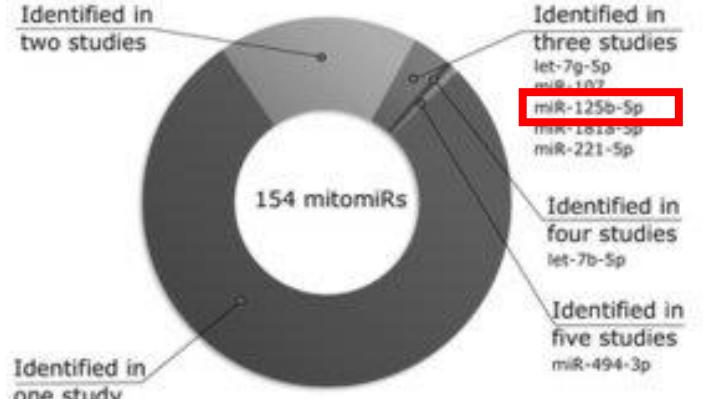
Research Article

Interplay of mitochondrial metabolism and microRNAs

Julian Geiger¹ • Louise T. Dalgaard¹

a

Co-occurrence of detected mitomiRs in published studies



miR-15a/miR-16 induces mitochondrial dependent apoptosis in breast cancer cells by suppressing oncogene BMI1

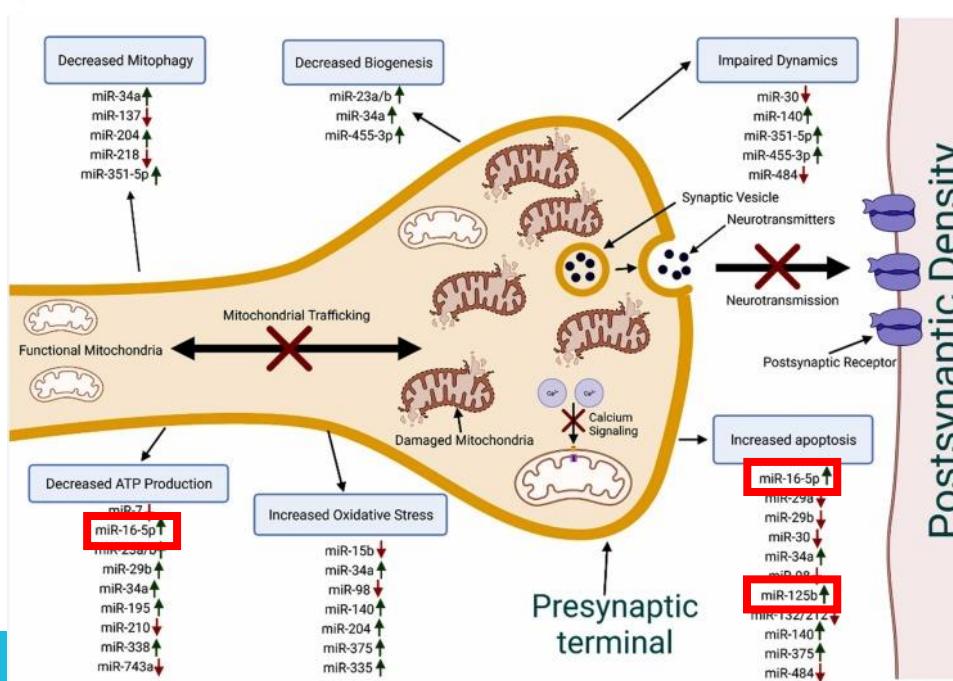
Nibedita Patel ^{a,1}, Koteswara Rao Garikapati ^{a,1}, M. Janaki Ramaiah ^{a, b}, Kavi Kishor Polavarapu ^d, Utpal Bhadra ^c, Manika Pal Bhadra ^a

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<https://doi.org/10.1016/j.lfs.2016.08.028>

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Combinatorial Effect of DCA and Let-7a on Triple-Negative MDA-MB-231 Cells: A Metabolic Approach of Treatment

Praveen Sharma, MSc¹ and Sandeep Singh, PhD¹

Abstract

Dichloroacetate (DCA) is a metabolic modulator that inhibits pyruvate dehydrogenase activity and promotes the influx of pyruvate into the tricarboxylic acid cycle for complete oxidation of glucose. DCA stimulates oxidative phosphorylation (OXPHOS) more than glycolysis by altering the morphology of the mitochondria and supports mitochondrial apoptosis. As a consequence, DCA induces apoptosis in cancer cells and inhibits the proliferation of cancer cells. Recently, the role of miRNAs has been reported in regulating gene expression at the transcriptional level and also in reprogramming energy metabolism. In this article, we indicate that DCA treatment leads to the upregulation of let-7a expression, but DCA-induced cancer cell death is independent of let-7a. We observed that the combined effect of DCA and let-7a induces apoptosis, reduces reactive oxygen species generation and autophagy, and stimulates mitochondrial biogenesis. This was later accompanied by stimulation of OXPHOS in combined treatment and was thus involved in metabolic reprogramming of MDA-MB-231 cells.

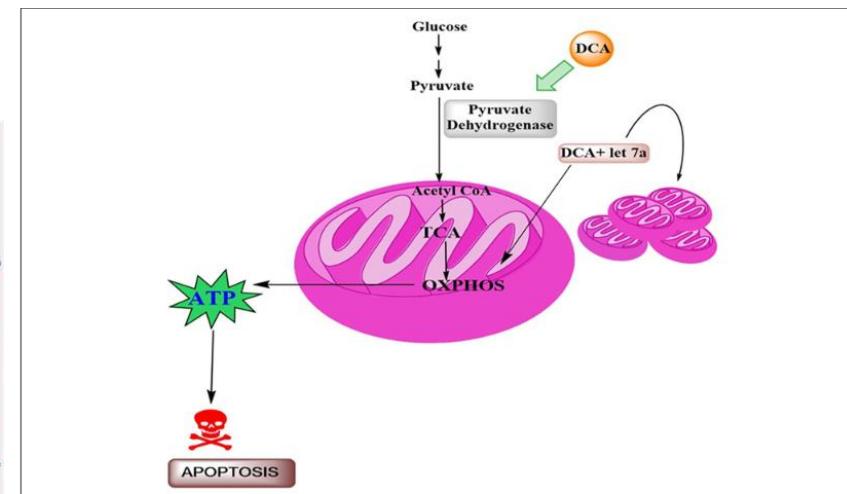


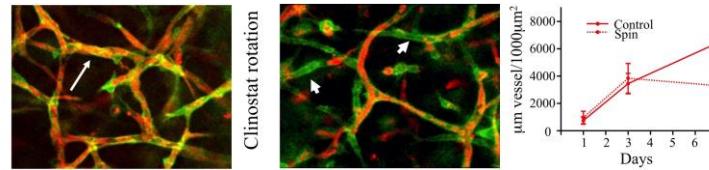
Figure 6. Model showing how combined treatment of let-7a and dichloroacetate (DCA) induces apoptosis by altering mitochondrial morphology and metabolism.

BMC Part of Springer Nature
Cancer Cell International
Home About Editors Submission Guidelines
Primary Research | Open Access | Published: 27 November 2021
Let-7a induces metabolic reprogramming in breast cancer cells via targeting mitochondrial encoded ND4
Praveen Sharma, Vibhu Sharma, Tarunver Singh Ahluwalia, Nilambara Dogra, Santosh Kumar & Sandeep Singh
Cancer Cell International 21 Article number: 629 (2021) | [Cite this article](#)
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Integrative Cancer Therapies
Volume 19: 1–11
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Targeted Based miRNA Therapy Mitigates Space Radiation Health Risks!!

PLOS ONE



- Utilized 3D human microvascular tissue models to determine functional impact of miRNAs and start development of miRNA based countermeasures.
- After 0.5Gy GCR irradiation Mature Vessels Collapse
- Applied combination of all 3 self delivery antagonists (AUM Biotech www.aumbiotech.com) **24 hours prior to irradiation** at 0.5μM for each antagonist, then cultured for 6 days after IR.

Simplified GCR Sim Irradiation



Peter Graham

COLUMBIA UNIVERSITY
IRVING MEDICAL CENTER

Ion species	Energy (MeV/n)	LET (keV/μm)	Dose (mGy)	Dose fraction (mGy)
Proton	1000	0.2	175	0.35
²⁸ Si	600	50.4	5	0.01
⁴ He	250	1.6	90	0.18
¹⁶ O	350	20.9	30	0.06
⁵⁶ Fe	600	173	5	0.01
Proton	250	0.4	195	0.39



AUMantagomir



Veenu Aishwarya
aum
LIFETECH

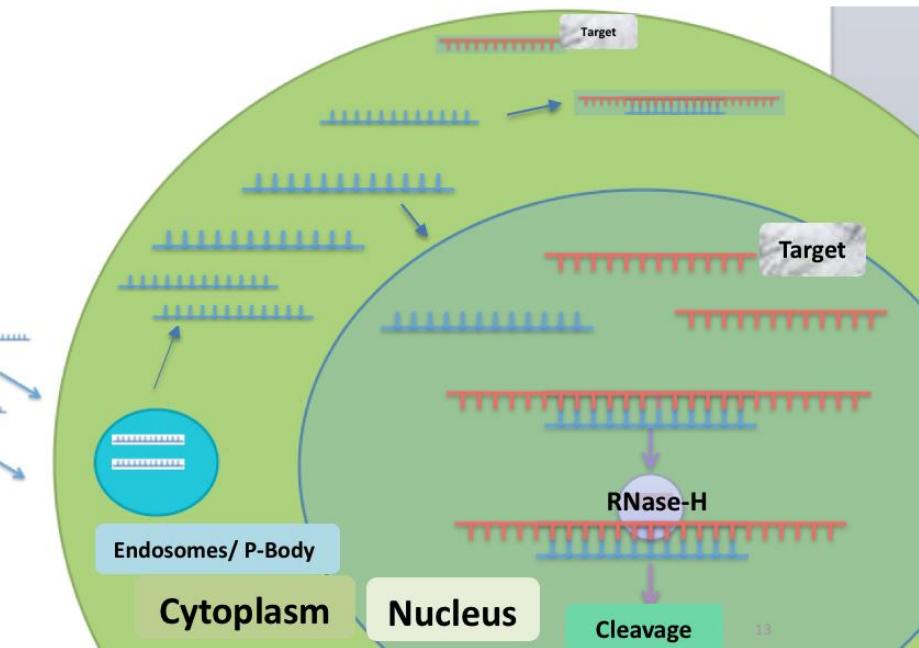
AUMsilence : 3rd Gen ASO Technology

Chemistry is the Key

- ✓ Self-delivery
- ✓ High Stability
- ✓ Can Target **Cytoplasmic & Nuclear RNA**
- ✓ High Potency
- ✓ High Sequence Specificity

Target:

- mRNA (AUMsilence)
- miRNA (AUMantagomir)
- lncRNA (AUMlnc)
- Viral RNA (AUMsilence V+)



miRNAs impact on Angiogenesis and Antagomir Countermeasures

LET-Dependent Low Dose and Synergistic Inhibition of Human Angiogenesis by Charged Particles: Validation of miRNAs that Drive Inhibition

Yen-Ruh Wuu • Burong Hu • Hazeem Okunola • ... Margareth Cheng-Campbell • Afshin Beheshti

Peter Graham Show all authors Show footnotes

Open Access • Published: November 25, 2020 • DOI: <https://doi.org/10.1016/j.isci.2020.101162>



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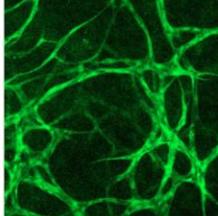


Highlights

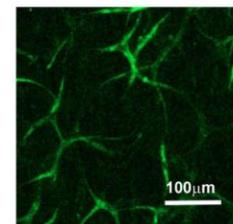
- Space radiation inhibits angiogenesis synergistically at low doses by 2 mechanisms
- Candidates for bystander transmission are microRNAs
- Three previously identified miRNAs showed downregulation of their angiogenesis targets
- Synthetic miRNA inhibitors were used to reverse the inhibition of angiogenesis



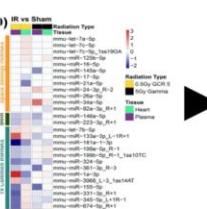
Human microvessels



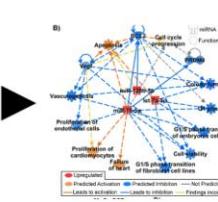
Space radiation



miRNA identification

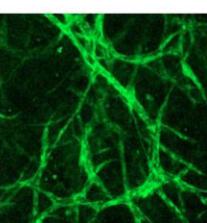


miRNA pathways



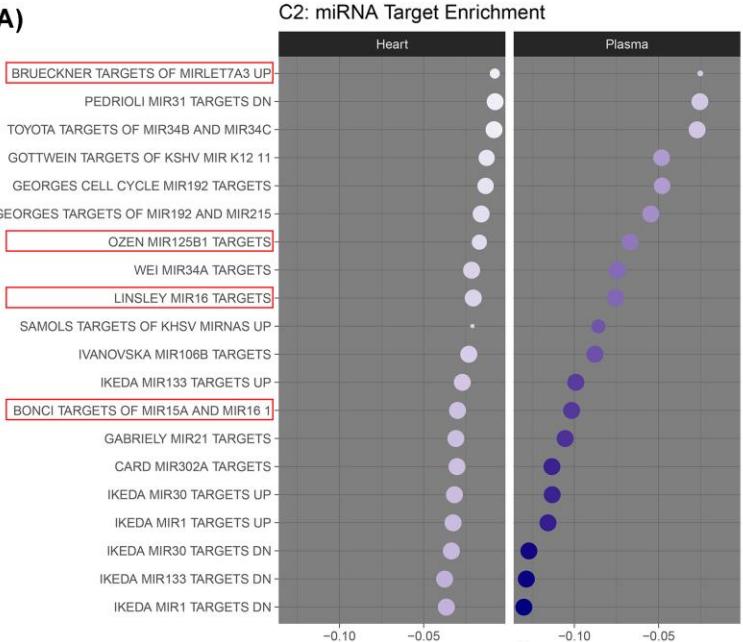
anti miR-16-5p
anti let-7a-5p
anti miR-125b-5p

miRNA inhibition

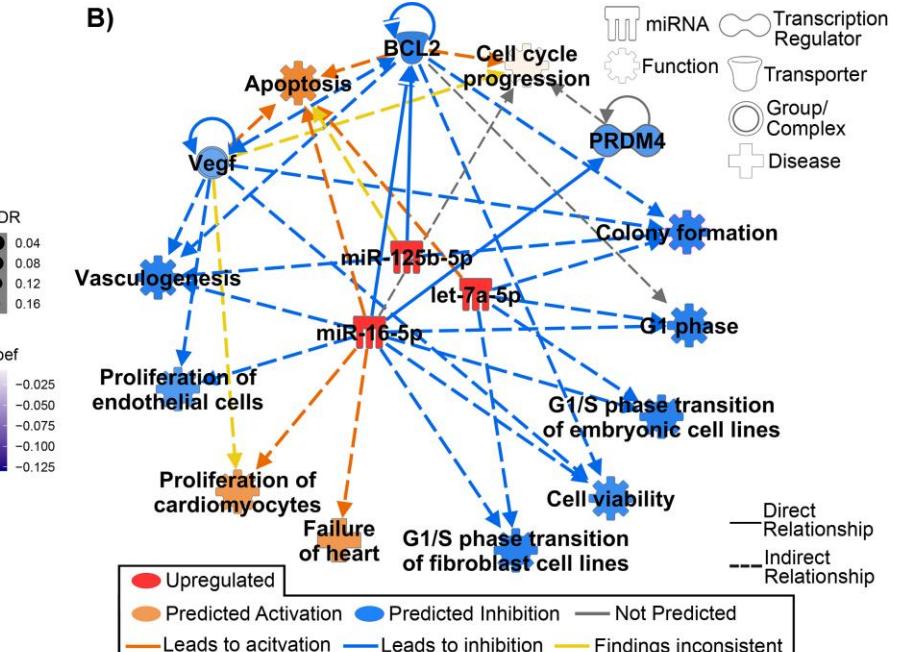


C2: miRNA Target Enrichment

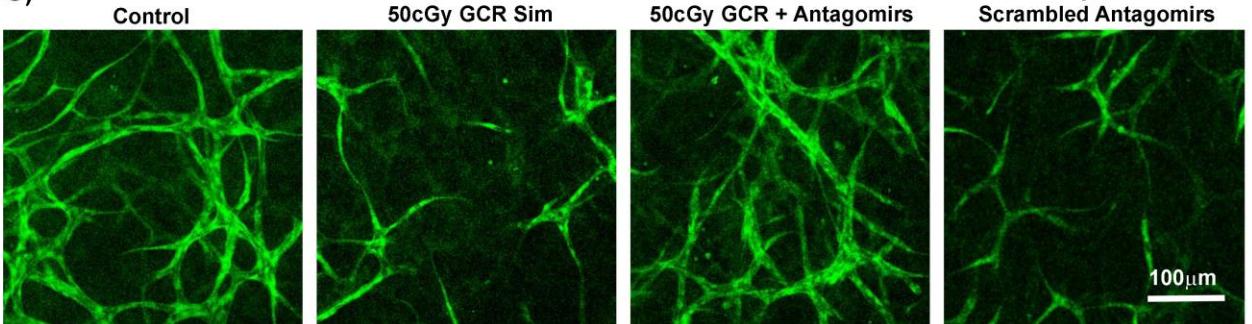
A)



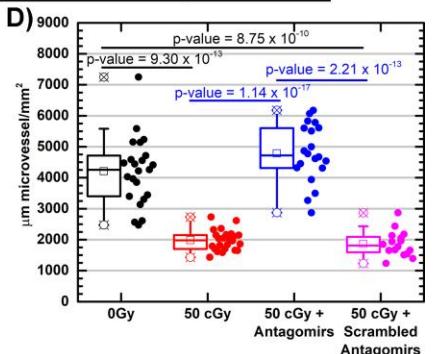
B)



C)



D)



Combination and Single Antagomir Treatment



Article PDF

So Long, and Thanks for All the Antagomirs: Space Radiation Damage Rescued by Inhibition of Key Spaceflight Associated miRNAs

J. Tyson McDonald, Lily Farmerie, Meghan Johnson, Jiwoon Park, and 19 more

This is a preprint; it has not been peer reviewed by a journal.

<https://doi.org/10.21203/rs.3.rs-2370597/v1>
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Status: Under Review



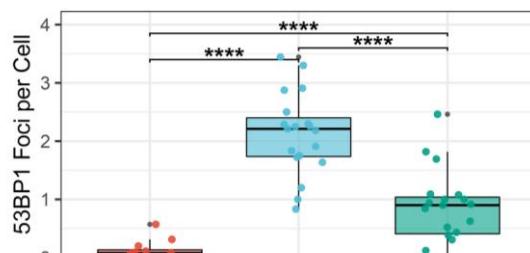
nature portfolio

Version 1
posted 09 Jan, 2023



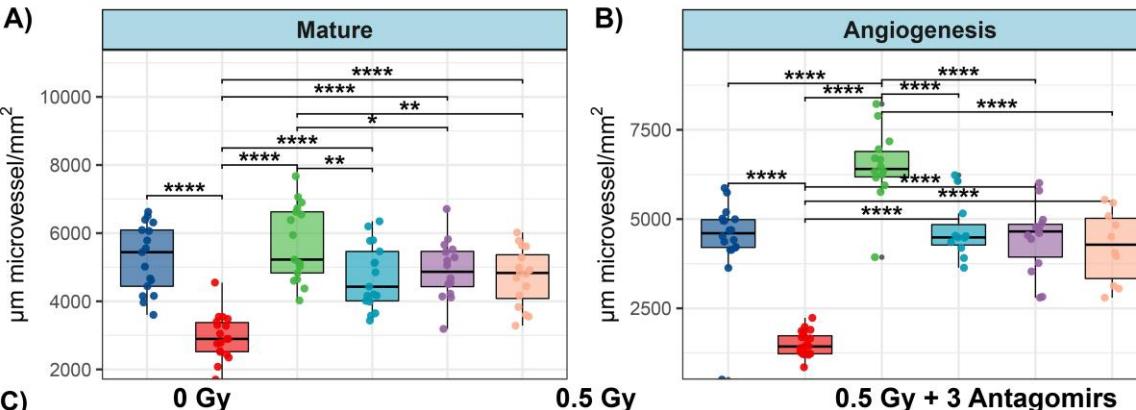
Antagomir Treatment on 3D Microvascular Tissues after GCR

Condition ■ 0.0 Gy ■ 0.5 Gy ■ 0.5 Gy + antagonists

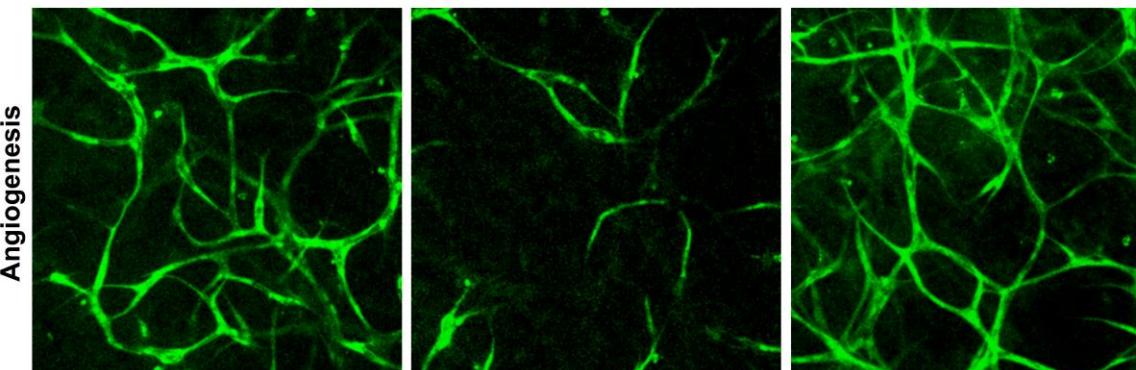
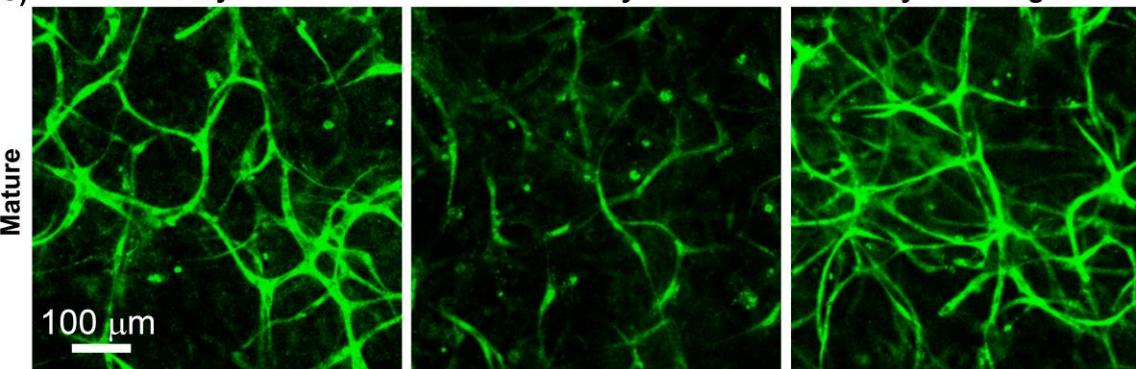


Antagomir Treatment on 3D Microvascular Tissues after GCR

Condition ■ 0.0 Gy ■ 0.5 Gy + 3 antagonists ■ 0.5 Gy + miR-16-5p antagomir
■ 0.5 Gy ■ 0.5 Gy + let-7a-5p antagomir ■ 0.5 Gy + miR-125b-5p antagomir



* p-value < 0.05
** p-value < 0.01
*** p-value < 0.001
**** p-value < 0.0001

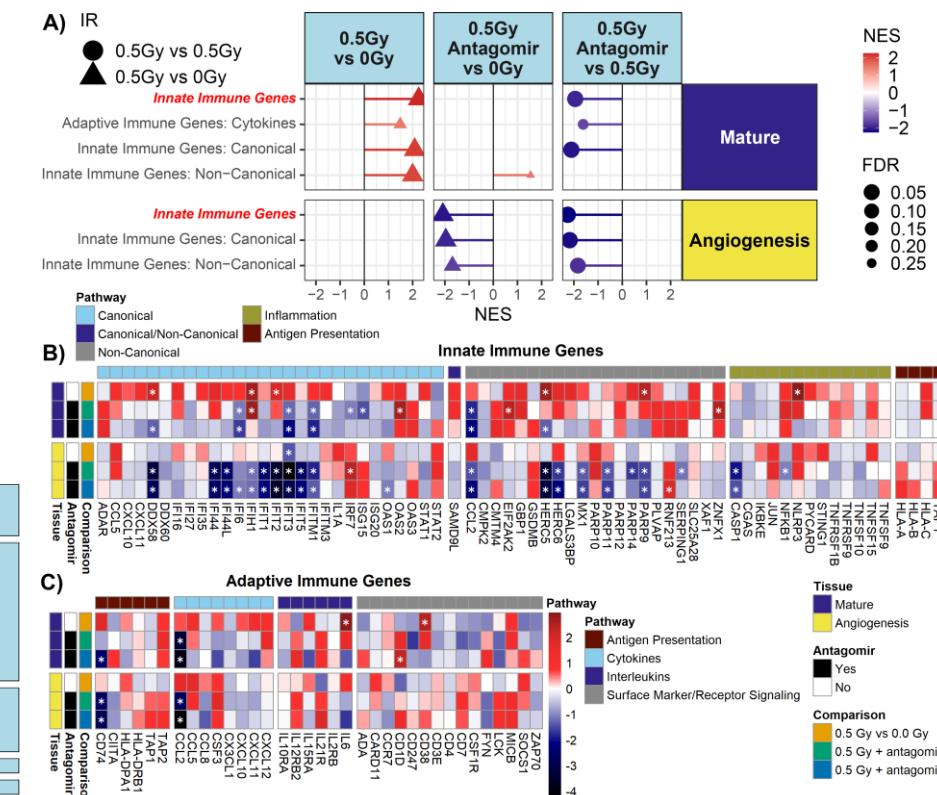
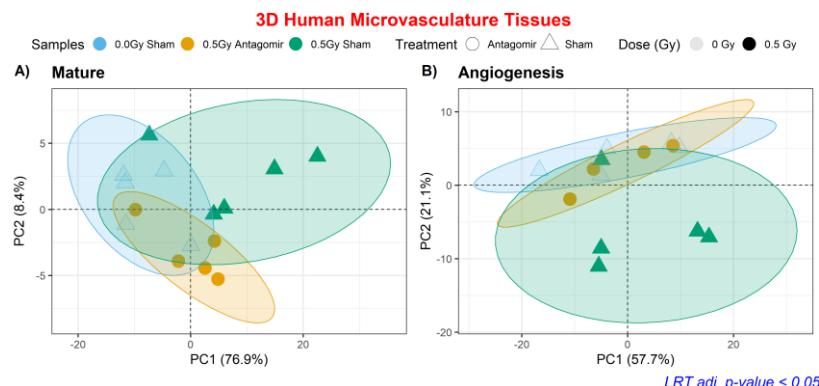
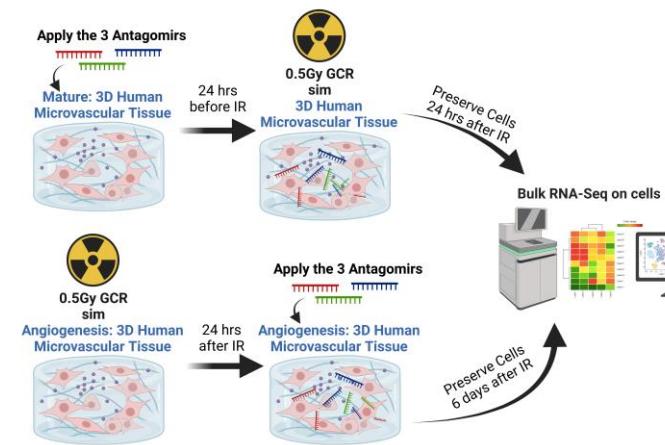


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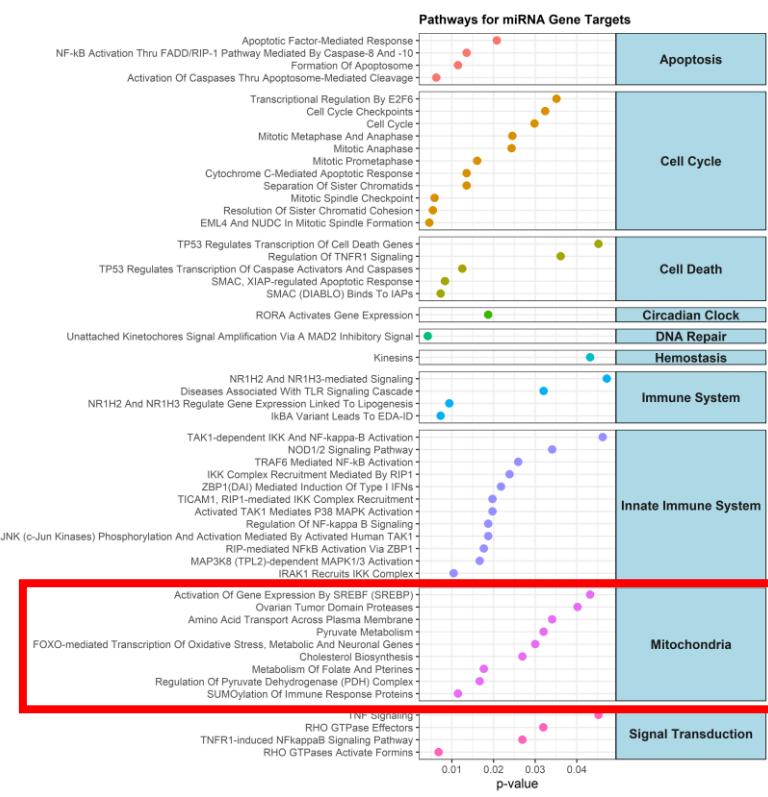
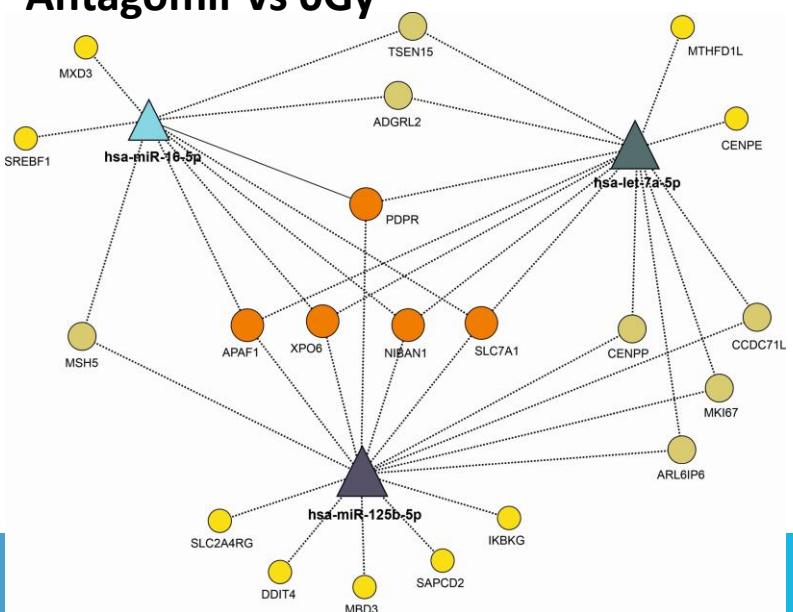
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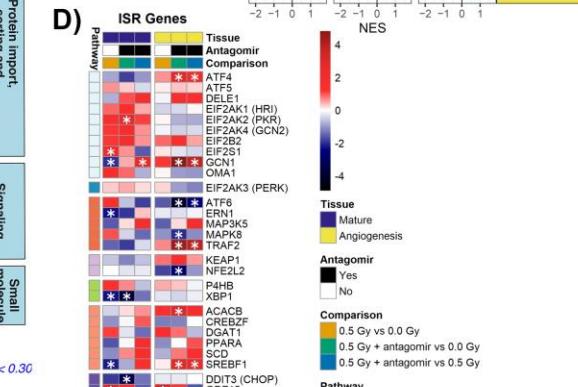
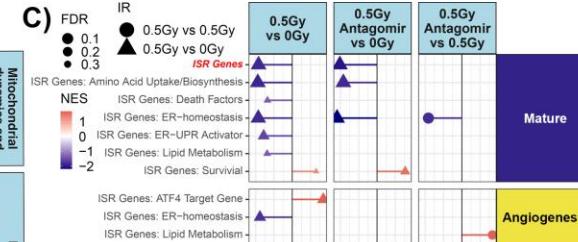
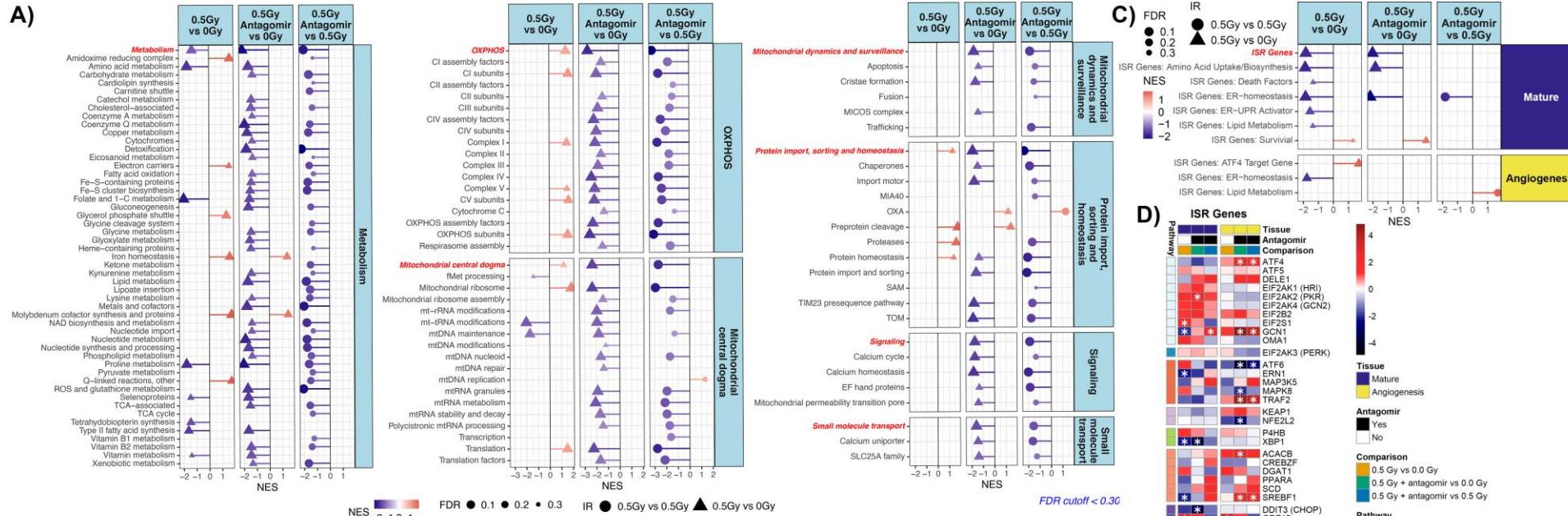
In vitro experiments with antagonists: Further analysis with the impact of antagonists on the 3D tissue model



Significantly Regulated Gene targets for all 3 miRNAs with 0.5Gy vs 0Gy, but are not significant anymore for 0.5Gy
Antagomir vs 0Gy



Further analysis with the impact of antagonirs on the 3D tissue Mature model: Mitochondrial Impact!



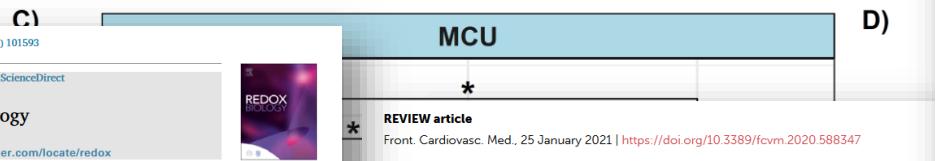
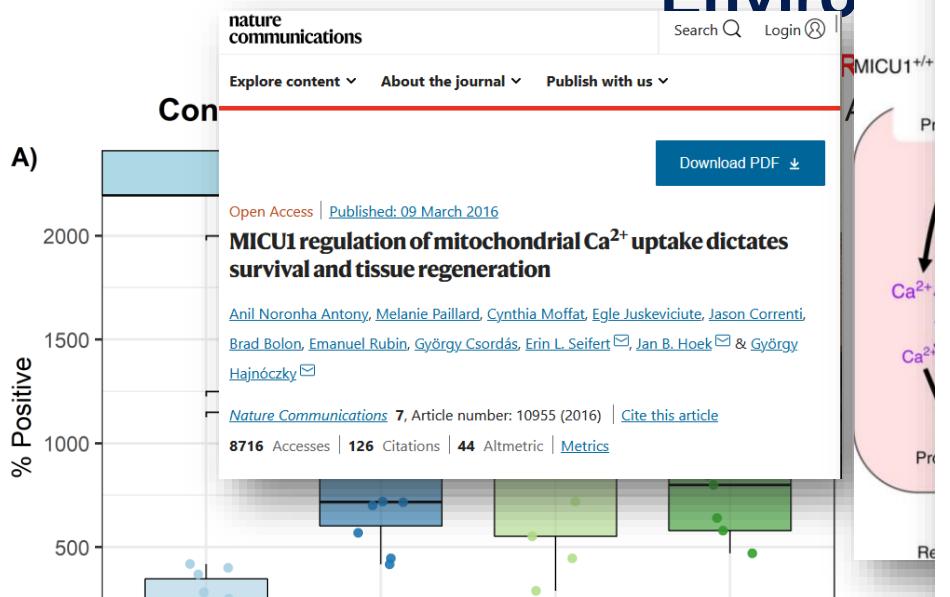
In vivo experiments with antagonists: Cardio Enviro



Peter Jirak



Paracelsus Medical
University, Salzburg



TGF- β 1 attenuates mitochondrial bioenergetics in pulmonary arterial endothelial cells via the disruption of carnitine homeostasis

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^aCenter for Blood Disorders, Medical College of Georgia at Augusta University, Augusta, GA, 30912, USA

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Keywords:
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TGF- β 1
PPAR
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NADPH oxidase
Mechanical forces

ABSTRACT

Transforming growth factor beta-1 (TGF- β 1) signaling is increased and mitochondrial function is decreased in multiple models of pulmonary hypertension (PH) including lambs with increased pulmonary blood flow (PBF) and pressure (Shunt). However, the potential link between TGF- β 1 and the loss of mitochondrial function has not been investigated and was the focus of our investigations. Our data indicate that exposure of pulmonary arterial endothelial cells (PAEC) to TGF- β 1 disrupted mitochondrial function as determined by enhanced mitochondrial ROS generation, decreased mitochondrial membrane potential, and disrupted mitochondrial bioenergetics. These events resulted in a decrease in cellular ATP levels, decreased hsp90/NOS interactions and attenuated shear-mediated NO release. TGF- β 1 induced mitochondrial dysfunction was linked to a nitration-mediated activation of Akt1 and the subsequent mitochondrial translocation of endothelial NO synthase (eNOS) resulting in the nitration of carnitine acetyl transferase (CAT) and the disruption of carnitine homeostasis. The increase in Akt1 nitration correlated with increased NADPH oxidase activity associated with increased levels of p47^{phos}, p67^{phos}, and Rac1. The increase in NADPH oxidase was associated with a decrease in peroxisome proliferator-activated receptor gamma (PPAR γ) expression. The presence of eNOS nitrated the primary substrate carnitine, thus disrupting the disruptive effect of TGF- β 1 on mitochondrial bioenergetics. Together, our studies reveal for the first time, that TGF- β 1 can disrupt mitochondrial function through the disruption of cellular carnitine homeostasis and suggest that stimulating carnitine homeostasis may be an avenue to treat pulmonary vascular disease.

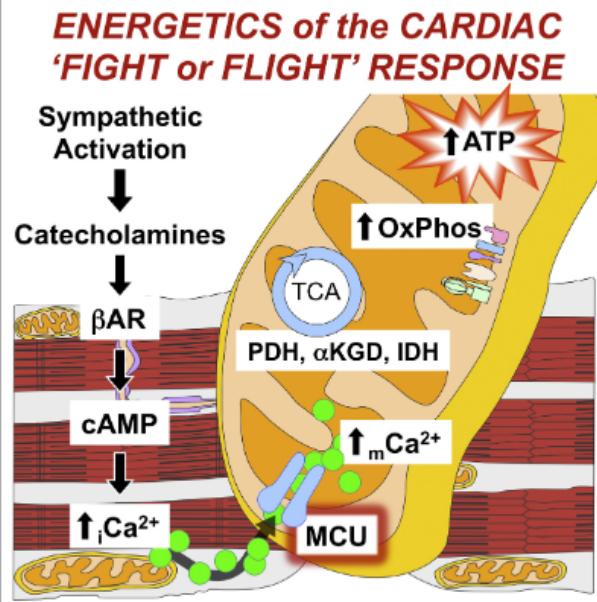
Cell Reports

Cell Death
& Disease

Search Q Report

The Mitochondrial Calcium Uniporter Matches Energetic Supply with Cardiac Workload during Stress and Modulates Permeability Transition

Graphical Abstract



Authors

Timothy S. Luongo, Jonathan P. Lambert, Ancai Yuan, ..., Joseph Y. Cheung, Muniswamy Madesh, John W. Elrod

Correspondence

elrod@temple.edu

In Brief

Luongo et al. show, using a conditional knockout mouse model, that the mitochondrial Ca²⁺ uniporter (MCU), although dispensable for homeostatic function, is necessary for the cardiac "fight-or-flight" contractile response and a significant contributor to mitochondrial permeability transition during ischemia-reperfusion injury.

Highlights

- The MCU is dispensable for baseline homeostatic cardiac function
- Deletion of *Mcu* protects against myocardial IR injury by reducing MPTP activation
- The MCU is required to match energetics with contractile demand during stress
- A slow MCU-independent uptake mechanism may maintain basal matrix mCa²⁺ content

Luongo et al., 2015, Cell Reports 12, 23–34
July 7, 2015 ©2015 The Authors
<http://dx.doi.org/10.1016/j.celrep.2015.06.017>

Fibrosis Gene Expression

CellPress

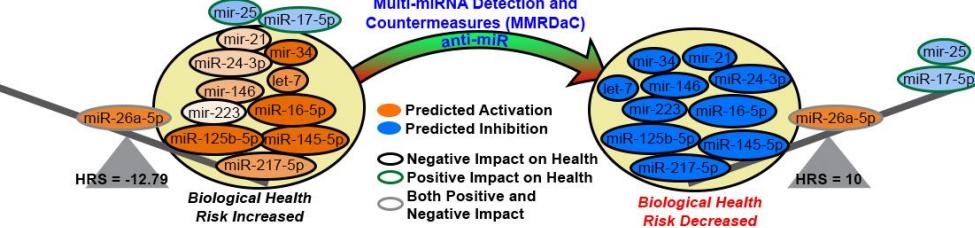
Summary of the Work



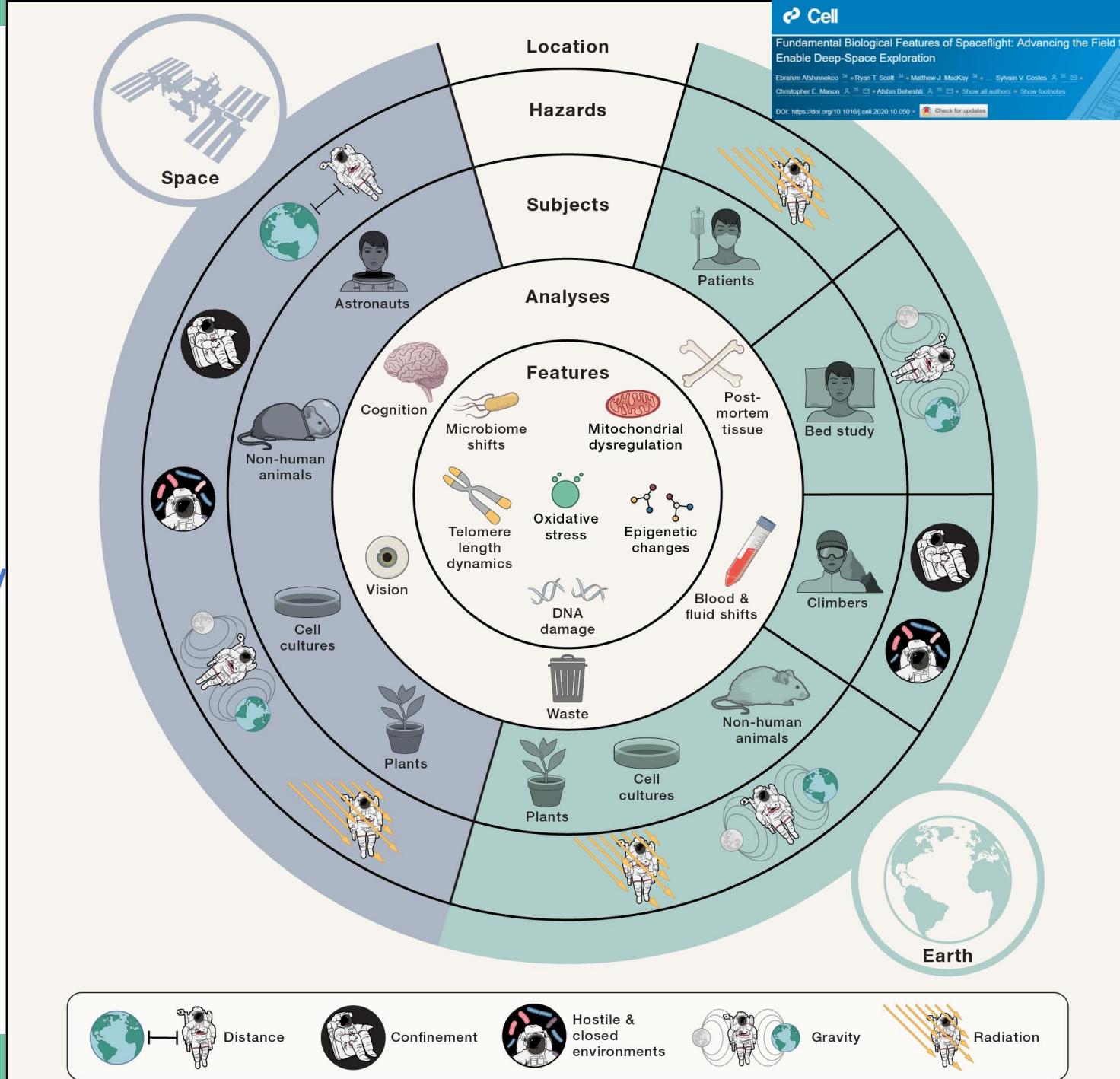
Are you calling me dysfunctional?



Dysregulates other key biological processes



- This spaceflight associated miRNA signature can be a novel minimally invasive biomarker to monitor increased health risks for long-term space missions.
- Also can be used for development of novel miRNA based therapeutic/countermeasure



Mitochondrial Paper Work Acknowledgements



Willian da Silveira



Deanne Taylor



Hossein Fazelinia



Komal Rathi



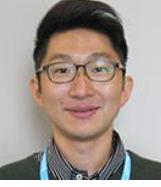
Douglas Wallace



Larry Singh



Benjamin
Stear



"Jimmy"
Mansuck Kim



Kathleen Fisch



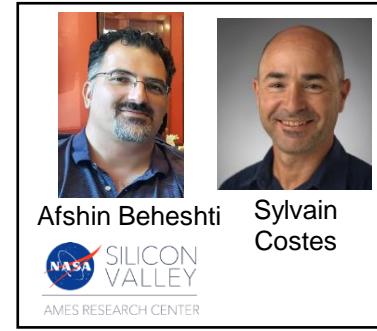
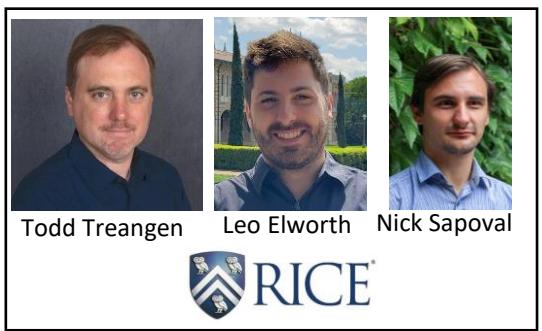
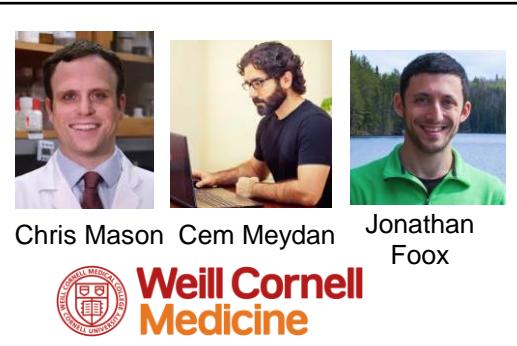
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Elizabeth Blaber



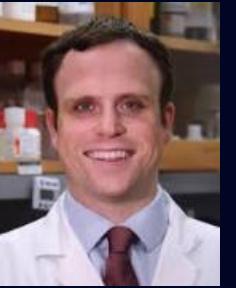
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Haltom



Rensselaer



Charles
Vanderburg



Diego Galeano



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**Appendix G: Solicitation of
Proposals for Flight and
Ground Space Biology
Research**



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